10, Appl 8, Appli 1, Appli

Sequence

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Sequence 1, Appli
Sequence 6, Appli
Sequence 6, Appli
Sequence 10, Appli
Sequence 10, Appli
Sequence 14, Appli
Sequence 2, Appli
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Sequence 119, Appli
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Pred. No. 7.8e-30;
0; Mismatches 209;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTESO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/824,873
FILING DATE: Filed Herewith
US-09-156-807-1

US-08-948-616-6

US-09-193-510-6

US-09-368-402-6

US-09-161-015-1

US-09-187-341-150

US-08-232-463-14

US-08-246-551-4

US-08-76-551-4

US-08-76-551-4

US-08-76-551-4

US-08-76-551-4

US-09-15-602-2

US-09-15-602-2

US-09-15-22-1

US-09-18-23-10

US-09-397-787-119

PCT-USS6-121298-1

US-09-397-787-119

US-09-397-787-119

US-09-397-787-119
                                                                                                                                                                                                                                                                            US-09-801-052-1
                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Guegler, Karl
TITLE OF INVENTION: NOVEL RAB PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                             ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PF-0240 US
                                                                                                                                                                                                                                                                                                                                                                                         US-08-824-873-2; Sequence 2, Application US/08824873; Patent No. 5843717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 1340 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLLOGY: linear
IMMEDIATE SOURCE:
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Best Local Similarity 59.0

Matches 305; Conservative
 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LIBRARY: PANCNOT04
CLONE: 738957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Palo Alto
STATE: CA
 USA
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Sequence 4, Appli
Sequence 10, Appl
Sequence 142, App
Sequence 6, Appli
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68, Appii
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33, Appii
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12643.623 Million cell updates/sec
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                                                                                                             January 16, 2003, 02:39:24; Search time 79 Seconds
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
                GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd
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US-09-198-1184-2

US-08-916-901-4

US-09-075-454-10

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US-08-773-423-4

US-08-399-913-66

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US-08-741-411-4
US-08-765-551-6
US-09-075-454-11
US-09-075-454-11
US-09-149-111-2
US-09-149-1411-2
US-09-149-14-11
US-09-165-979-1
US-09-184-13
US-09-184-14-16
US-09-18-14-16
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US-08-247-946A-5
PCT-US95-06420-5
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Maximum Match 100%
Listing first 45 summaries
                                                                              - nucleic search, using sw model
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Maximum DB seq length: 2000000000
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Match
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85.8
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                22 TACGACGTCGCTTCAAGGTCATGCTGGTGGGGGACTCGGGTGTGGGGAAGACCTGTG
                                                       117 CTGTGC-CGCTTCACCGACAACGAGTTCCACTCCTCGCACATCTCCACCATCGGTG--TT
                                                                        174 GACTITAAGATGAAGACCATAGAGGTAGACGGCATCAAAGTGCGGATACAGATCTGGGAC
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APPLICATION NUMBER: US/09/198,184 FILLING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Incyte Pharmaceuticals, Inc. STREET: 3174 Porter Drive CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/09198184
Patent No. 6010859
GENERAL INFORMATION
APPLICANT: Hillman, Jennifer L.
TITLE OF INVENTION: NOVEL RAB PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: PF-0240 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/824,873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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                                                                                                                                                                                          Length 1340;
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                                                                                                                                                                                          Score 156.6; DB 3;
Pred. No. 7.8e-30;
0; Mismatches 209;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Hillman, Jennifer L. APPLICANT: Lal, Preeti APPLICANT: Corley, Neil C. APPLICANT: Shah, Purvi TITLE OF INVENTION: RAB PROTEINS NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-916-901-4; Sequence 4, Application US/08916901; Sequence No. 5892012; Patent No. 592012; GENERAL INFORMATION:
                                                                                                                                                                                            4.8%;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                LENGTH: 1340 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                     Matches 305; Conservative
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Ph
                                                                                         IMMEDIATE SOURCE:
LIBRARY: PANCNOT04
CLONE: 738957
                                                                            linear
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                                                                                                                                                                                                                   Best Local Similarity
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                                                                            TOPOLOGY:
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                                                                                                                                                     US-09-198-184-2
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                                                                                                                                                                                              Query Match
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APPLICANT: Corley, Neil C.
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56.7%; Pred. No. 9.4e-30;
iive 0; Mismatches 220; Indels
SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                           PF-0367 US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/916,901 FILING DATE: Filed Herewith PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                   REFERENCE/DOCKET NUMBER:

TELECOMMUNICATION INFORMATION:

TELEFONE:

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TELEFONE:

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TENERATION FOR SEQ ID NO:

SEQUENCE CHARACTERISTICS:

LENGTH:

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TYPE:

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TYPE:

STRANDENNESS:

Single
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Patent No. 6300472
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                                                                                                                          NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749
                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                           Best_Local Similarity 56.7
Matches 288; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 6300472
GENERAL INFORMATION:
APPLICANT: Hillman, Jes
APPLICANT: Lal, Preeti
                                                                                                                                                                                                                                                                                                                               LIBRARY: LIVRTUTO4
CLONE: 2514506
                                                                                                                                                                                                                                                                                                   linear
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                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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US-08-916-901-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           297 TACCGGGGGGCTCATGGCATCATCGTGTGTATGACGTCACTGACCAGGAATCCTACGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  156 ATCTCCACCATCGGTGTTGACTTTAAGATGAAGACCATAGAGGTAGACGGCATCAAAGTG
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56.7%; Pred. No. 9.4e-30;
tive 0; Mismatches 220;
                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/154,602
FILING DATE:
                                                                                                        ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PF-0367 US
APPLICANT: Shah, Purvi
TITLE OF INVENTION: RAB PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/916,901
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                        IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 925 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMBIATE SOURCE:
LIBRARY: LIVRTUT04
CLONE: 2514506
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Matches 288; Conservative
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                                                                                                                                                                                                                                                                        Diskette
                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                           NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
                                                                                                                                                                                                   USA
                                                                                                                                                                                                                        94304
                                                                                                                                                                           STATE: CA
                                                                                                                                                                                                                                                                                        COMPUTER:
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                                                                                                                                                                                              COUNTRY:
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271 ACCGCTGGGCAGGAACGGTTCCGAAGCGTCACCCATGCTTATTACAGAGATGCTCAGGCC 330
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                                                                                                                                                                                                                                                                                                       ACTGCAGGCAGGAGAGATACCAGACCATCACAAAGCAGTACTATCGGCGGGCCCAGGGG
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174 GACTTTAAGATGAAGACCATAGAGGTAGACGCCATCAAAGTGCGGATACAGATCTGGGAC
                                                                                         234 ACTGCAGGCAGGAGAGATACCAGACCATCACAAAGCAGTACTATCGGCGGGCCCAGGGG
                                                                                                                                             294 ATATTTTTGGTCTATGACATTAGCAGCGAGCGCTCTTACCAGCACATCATGAAGTGGGTC
                                                                                                                                                                                                                  354 AGTGACGTGGATGAGTACGCACCAGAAGGCGTCCAGAAGATCCTTATTGGGAATAAGGCT
                                                                                                                                                                                                                                                                                      414 GATGAGGAGCAGAAACGGCAGGTGGGAAGAAGAGGAAGGGCAGCAGCTGGCGAAGGAGTAT
                                 192 GACTICAGGAACAAGGIGGIGACIGIGGAIGGCGIGAGAGIGAAGCIGCAGAICIGGGAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. 6426186 412477.1CB1
US-09-484-9708-142
                                                                                                                                                                                                                                                                                                                                                                              GGCATGGACTTCTATGAAACAAGTGCCTGCACCAACCTCAA 514
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Jones, Karen A.
APPLICANT: Volkmuth, Mayne
APPLICANT: Walker, Michael G.
TITLE OF INVENTION: BONE REMODELING GENES
FILE REFERENCE: PB-0014 US
CURRENT APPLICATION NUMBER: US/09/484,970B
CURRENT FILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 172
SOFTWARE: PERL PLOGITAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 142, Application US/09484970B
; Patent No. 6426186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 4.29
Best Local Similarity 57.55
Matches 265; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Jones, 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-484-970B-142
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LENGTH: 2612
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Pred. No. 8.6e-25;
0; Mismatches 193;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PF-0168-1 CIP
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                                                         516 ATTAAAGAGTCATTCACGCGTCTGACAG 543
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                                                                                           537 GICGAGCAGGCGITCAIGACCAIGGCIG 564
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Word Perfect 6.1/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/075,454
FILING DATE: Herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA
APPLICATION NUBER: 08/766,551
FILING DATE: DECEMBER 12, 1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                Sequence 10, Application US/09075454 Patent No. 6391580
                                                                                                                                                                                                                                                                                                                                                                                RAS PROTEINS
                                                                                                                                                                                                                                                   APPLICANT: Tang, Y. Tom
APPLICANT: Lal, Preeti
APPLICANT: Call, Preeti
APPLICANT: Guegler, Karl J.
APPLICANT: Corley, Neil C.
APPLICANT: Patterson, Chandra
APPLICANT: Batta, Sajeev
APPLICANT: Baughn, Mariah R.
TITLE OF INVENTION: RAS PROTEIN
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                        Hillman, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Cerrone, Michael C.
REGISTRATION NUMBER: 39,132
REFERENCE/DOCKET NUMBER: PF-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1: Diskette
IBM Compatible
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LENGTH: 875 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best_Local Similarity 57.5
Matches 265; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: UCMCL5T01
CLONE: 1528559
                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                       INFORMATION:
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262 TCACAAAGCAGTACTATCGGCGGGCCCAGGGGATATTTTGGTCTATGACATTAGCAGCG 321
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                                                                                                                                                                                                                                                                          420 CGATCGTCGTCATGCTCGTGGGTAACAAAGTGACCTCAGCCAGGGCCGGGAAGTGCCCA 479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           51 AAGCAGTACGATGTGCTGTTCCGGCTGCTGTCGGGGACTCCGGGGTGGGCAAGACC 110
  322 AGCGCTCTTACCAGCACATCATGAAGTGGGTCAGTGACGTGGATGAGTACGCACCAGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                             480 CTGAGGGCCCGAATGTTCGCTGAAACAATGGACTGCTCTTCCTGGAGACCTCAGCCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Rhodes, Kenneth
APPLICANT: Betty, Maria
APPLICANT: Ling, Hual Ping
APPLICANT: Ling, Hual Ping
APPLICANT: An, Wengian
TITLE OF INVENTION: POTASSIUM CHANNEL INTERACTORS AND USES THEREFOR
FILE REFERENCE: MNI-070622
                                                                    202 ACGGCATCAAAGTGCGGATACAGATCTGGGACACTGCAGGGCAGGAGAGATACCAGACCA
                                                                                                              240 GCACCGCTGCTGTCAAGGCTCAGATCTGGGACACAGCTGGCCTGGAGCGGTACCGAGCCA
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52.2%; Pred. No. 5.1e-19;
Live 0; Mismatches 230;
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CURRENT FILING DATE: 1999-09-21

BARLIER APPLICATION NUMBER: USN 60/110,277

BARLIER FILING DATE: 1998-11-30

BARLIER FILING DATE: 1998-11-25

BARLIER FILING DATE: 1998-11-25

BARLIER FILING DATE: 1998-11-25

BARLIER FILING DATE: 1998-11-25

BARLIER FILING DATE: 1998-11-26

BARLIER APPLICATION NUMBER: USN 60/109,333

BARLIER FILING DATE: 1999-04-23

BARLIER FILING DATE: 1999-04-23

BARLIER FILING DATE: 1999-07-09

BARLIER FILING DATE: 1999-07-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 66, Application US/09399913 Patent No. 6361971
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Best Local Similarity 52.2
Matches 251; Conservative
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US-09-399-913-66
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TGATCGGGGACTCCGGGGTGGGCAAGACCTGCTGTGCCGCTTCACCGACAACGAGT 141
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                                                                GATGAGGAGCAGAAACGGCAGGTGGGAAGAGAGCAAGGGCAGCAGCTGGCCGAAGGAGTAT
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                                                                                                                                                     GGCATGGACTTCTATGAAACAAGTGCCTGCACCAACCTCAA 514
                                                                                                                                                                              3.7%; Score 120.6; DB 2;
51.5%; Pred. No. 7.1e-21;
tive 0; Mismatches 280;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Goli, Surya K.

APPLICANT: Bandman, Olga
TITLE OF INVENTION: NOVEL RAB PROTEINS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PF-0183 US
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FILING DATE: Herewith
                                                                                                                                                                                                                                                                                                          Sequence 4, Application US/08773423
Patent No. 5869291
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Hillman, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 847 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 301; Conservative
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: Consensu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   230
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               9
                                                                                                                                171 GTTGACTTTAAGATGAAGACCATAGAGGTAGACGGCATCAAAGTGCGGATACAGATCTGG
                                                                                                                                                                          231 GACACTGCAGGGCAGGAGATACCAGACCATCACAAAGCAGTACTATCGGCGGGCCCAG
                                                                                                                                                                                                      181 GATACAGCAGGCAGGAGTCCTTTCGTTCTATCACAAGGTCATATTACAGAGGTGCAGCG
                                                                                                                                                                                                                                       GGGATATTTTGGTCTATGACATTAGCAGCGAGCGCTCTTACCAGCACATCATGAAGTGG
                                                .11 TGCCTGCTGTGCCGCTTCACCGACAACGAGTTCCACTCCTCGCACATCTCCACCATCGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED TITLE OF INVENTION: TO ALZHEIMER'S DISEASE AND USES THEREFOR NUMBER OF SEQUENCES: 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ν
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: LERNER, DAVID, LITTENBERG, KRUMHOLZ & MENTLIK
STREET: 600 SOUTH AVENUE WEST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: SCHERING 3.0-017 CIP CIP TELECOMMUNICATION INFORMATION:
TELEPHONE: (908) 654-5000
TELEFAX: (908) 654-5066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING STATE SOFTWARE: ASCII CURRENT APPLICATION NUMBER: US/08/888,077A FILING DATE: 03-UUL-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 28, Application US/08888077A
Patent No. 6020143
GENERAL INFORMATION:
APPLICANT: ST. GEORGE-HYSLOP, PETER
APPLICANT: ROMMENS, JOHANNA M
APPLICANT: FRASER, PAUL E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/592,541
FILING DATE: 26-JAN-1996
ATTONEY/AGENT INFORMATION:
NAME: PALISI, THOMAS M
REGISTRATION NUMBER: 36,629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                       TGCTGTGCCGCTTCACCGACAACGAGTTCCACTCCTCGCACATCTCCACCATCGGTGTTG 174
                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTGCAGGGCAGGAGATACCAGACCATCACAAAGCAGTACTATCGGCGGGCCCAGGGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       373 AAGAACTGAGAGATCATGCTGATAGTAACATTGTTATCATGCTTGTGGGCAATAAGAGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     433 ATCTACGTCATCTCAGGGCAGTTCCTACAGATGAAGCAAGAGCTTTTGCAGAAAAGAATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACTITAAGATGAAGACCATAGAGGTAGACGCCATCAAAGTGCGGATACAGATCTGGGACA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       355 GTGACGTGGATGAGTACGCACCAGAAGGCGTCCAGAAGATCCTTATTGGGAATAAGGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             295 TATTTTTGGTCTATGACATTAGCAGCGAGCGCTCTTACCAGCACATCATGAAGTGGGGTCA
                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                                                           Length 970
                                                                                                                                                                                                                       Score 110; DB 3; Length 97
Pred. No. 3.5e-18;
0; Mismatches 210; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Bandman, Olga
APPLICANT: Au Toung, Janice
TITLE OF INVENTION: NOVEL RAB PROTEINS
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: INCYTE PHARMACEUTICALS, STREET: 3174 Porter Drive CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US/08/741,411
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     493 GTTTGTCATTCATTGAAACTTCGGCC 518
                                                                                                                       ; NAME/KEY: misc_feature
; LOCATION: 1..970
; OTHER INFORMATION: /note= "Y2H9"
US-08-888-077A-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/741.4
FILING DATE: Herewitt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-741-411-6; Sequence 6, Application US/08741411; Patent No. 6124116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                             3.4%;
                                                                                                                                                                                                                             Query Match 3.45
Best Local Similarity 52.95
Matches 236; Conservative
                                     : 970 base pairs
nucleic acid
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                           single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: CA
COUNTRY: US
ZIP: 94304
                                                                       STRANDEDNESS
                                                                                             TOPOLOGY:
                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                   115
                                                                                                                                                                                                                                                                                                                                                                                                                                                             175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       235
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1800 Diagonal Road, Suite 500
                    Alexandria
                                                                                                                                                                                                FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CCAGGGGATATTTTGGTCTATGACATTAGCAGCGAGCGCTCTTACCAGCACATCATGAA 346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GTGGGTCAGTGACGTGGATGAGTA----CGCACCAGAAGGCGTCCAGAAGATCCTTATTGG 403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  404 GAATAAGGCTGATGAGGAGCAGAAACGGCAGGTGGGAAGAGAGCAAGGGCAGCAGCTGGC 463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                464 GAAGGAGTATGGCATGGACTTCTATGAAACAAGTGCCTGCACCAACCTCAACATTAAAGA 523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              518 CGACAGCCAGAAGTTGCTGTTCATGGAAACTTCGGCCAAACTGAACCACCAGGTGTCGGA 577
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      167 CGGTGTTGACTTTAAGATGAAGACCATAGAGGTAGACGCCATCAAAGTGCGGATACAGAT 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     218 CGTGGGTGCGTTCTTCACAAAGGAGGTGGATGTGGGGTGCCACCTCTGAAGCTTGAGAT 277
                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 820;
                                                                                                                                                                                                                                                                                                                                                                                      Score 91.2; DB 3; Length 8
Pred. No. 1.7e-13;
0; Mismatches 203; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: DORNER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: SCHEIFLINGER, F.
APPLICANT: FALKNER, F. G.
TITLE OF INVENTION: RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY & LAIGHER
                                                                                                                PF-0139 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14, Application US/08232463
Patent No. 5670367
                                                                            NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-
TELECOMMUNICATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                        2.8%;
                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 820 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 226; Conservative
                                                                                                                                              TELEPHONE: 415-855-05: TELEFAX: 415-845-4166
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            584 CCGGATGCGTGC 595
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US-08-741-411-6
                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                           TYPE: nucleic
STRANDEDNESS:
TOPOLOGY: line
                                                                                                                                                                                                                                                                                                         IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                              FILING DATE:
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                                                                                                                                                                                                                                                                                                                           LIBRARY:
                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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984 CAGCCAGGGCTGGAGTCCAGGCCACTTTCAGCTGCTCCTTTCTCCGTGCATCGTGTCTCT 1043
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                864 TGTCCCCAGCCCGTCCACCCTGGAGTGGTCTTCTTCAGCCTGTTTCCCCAGCCACAGGCC 923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           804 TAGCCACTACTCCCCCTGCCTGGCCTGAGAGCGGCTCTGCTGTCATCTCAAGCAGCCCC 863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 7218;
                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/232,463
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 85.8; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative 251; Mismatches 123;
                                                                                                                                                                                                                                                                              APPLICATION NUMBER: EP 91 114 300.6
AFTLING DAFE: 26-A0G-1991
ATTORNEY AGENT INFORMATION:
NAME: BENT, Stephen A.
RECISTRATION NUMBER: 29,768
REFERENCE/DOCKET NUMBER: 30472/114 IMMU
                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/935,313
FILING DATE:
                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         YYYYYYYYYYYYYYGTACCAAA 1442
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US-08-741-411-4
; Sequence 4, Application US/08741411
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEX: 899149
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (703)836-9300
TELEFAX: (703)683-4109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7218 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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Matches 9; Conserv
COUNTRY: USA
ZIP: 22313-0299
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Word Perfect 6.1/MS-DOS 6.2
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/075,454
FILING DATE: Herewith
                                                                                                                                                                                                                 APPLICANT: Baughn, Mariah R.
TITLE OF INVENTION: RAS PROTEINS
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/766,551
FILING DATE: DECEMBER 12, 1996
ATTORNEY/AGENT INFORMATION:
              US-09-075-454-8
; Sequence 8, Application US/09075454
; Patent No. 6391580
                                                      'atent No. occasion:
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
                                                                                                                                      APPLICANT: Guegler, Karl J.
APPLICANT: Corley, Neil C.
APPLICANT: Patterson, Chandra
APPLICANT: Batra, Sajeev
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
TELEFAX: 650-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.5%;
                                                                                                                                                                                                                                                                                                      3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
LENGTH: 1172 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 2.55
Best Local Similarity 54.4
Matches 161; Conservative
                                                                                                        Tang, Y. Tor
Lal, Preeti
                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LIBRARY: KIDNNOT05
CLONE: 627565
                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                           Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                              a
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                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-075-454-8
                                                                                                                                                                                                                                                                                                                                              STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96 GACCTGGGCGTGGGSAAGACCAGYATCATCAAGCGCTACGTCCACCACTTCTCCCAG 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                150 TCGCACATCTCCACCATCGGTGTTGACTTTAAGATGAAGACCATAGAGGTAGACGGCATC 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      276 CGAGTATACTACAAGGAAGCTGTTGGTGCTTTTGTAGTCTTTGATATATCAAGAAGTTCC 335
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36 CCGCCCCCAGGATGCAGGCCCCCCACAAGGACCACCTGTACAGTTGCTGGTGATTGC 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30 CCGCTGGCCCCAGTCATGGCGAAGCAGTACGATGTGCTGTTCCGGCTGCTGCTGATCGGG 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            267 AAGCAGTACTATCGGCGGGCCCAGGGGATATTTTTGGTCTATGACATTAGCAGCGGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 80.6; DB 3; Length 8
Pred. No. 8.2e-11;
2; Mismatches 151; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCTTACCAGCACATCATGAAGTGGGTCAGTGACGTGGAT 365
GENERAL INFORMATION:
APPLICANT: Bandman, Olga
APPLICANT: Au-Young, Janice
TITLE OF INVENTION: NOVEL RAB PROTEINS
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFRENCE/DOCKET NUMBER: PF-0139 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-845-055
TELEFAX: 415-845-4166
                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ VESION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/741,411
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.5%;
                                                                                                                         STREET: 3174 Porter Drive
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 890 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEC ID NO:
                                                                                                                                                                                                                                                                                                                                              Herewith
                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                      COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: cDNA IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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US-08-741-411-4
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                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                STATE:
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125 CTTCACCGACAACGAGTTCCACTCCTCGCACATCTCCACCATCGGTGTTGACTTTAAGAT 184 298 GGAACGATITCAGGIGCTGGGCATICCCTICAGITIGCAGCITIGGGATACCGCTGGGCA 357 245 GGAGAGATACCAGACCATCACAAGCAGTACTATCGGCGGGCCCAGGGGATATTTTGGT 304 238 GITCTGCAAAGACACCITTGAIAAGAAITACAAGGCCACCAITGGAGTGGACTTCGAGAT GAAGACCATAGAGGTAGACGGCATCAAAGTGCGGATACAGATCTGGGACACTGCAGGGCA DB 4; Length 1172; 0; Mismatches 135; Indels CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC. NAME: Cerrone, Michael C.
REGISTRATION NUMBER: 39,132
REFERENCE/DOCKET NUMBER: PF-0168-1 CIP

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us-09-817-198a-1.rni

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695 GGAGAGGTTCAAATGCATTGCATCAACCTACTATAGAGGAGCTCAAGCCATCATCATTGT 754
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CLONE: 1651593
US-09-075-454-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE: CA
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APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
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Matches 161;
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              305 CTATGACATTAGCAGCGCTCTTACCAGCACATCATGAAGTGGGTCAGTGACG 360
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Pred. No. 1.3e-10;
0; Mismatches 135; Indels
                                                                                                                                                                                             APPLICANT: Bandman, Olga
APPLICANT: Hawkins, Phillip R.
APPLICANT: Hawkins, Phillip R.
APPLICANT: Goll, Surya K.
APPLICANT: Goll, Surya K.
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PF-0168 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: DOS
SOFTWARE: FASTED Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/766,551
FILING DATE: Herewith
                                                                                                                              Sequence 6, Application US/08766551
Patent No. 5840569
GENERAL INFORMATION:
                                                                                                                                                                                   APPLICANT: Hillman, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEPAX: 415-845-4166
                                                                                                                                                                                                                                                                                                                     3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 2.5%;
Best Local Similarity 54.4%;
Matches 161; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                           IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: (
SEQUENCE CHARACTERISTICS:
LENGTH: 1255 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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CLONE: 627051
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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STATE: CA
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STREET: 31
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Pred. No. 1.5e-10;
0; Mismatches 135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/766,551

FILING DATE: DECEMBER 12, 1996

ATTORNEY/AGENT INFORMATION:

NAME: CETROME, MICHAEL C.

REGISTRATION NUMBER: 39,132

REFERENCE/DOCKET NUMBER: PF-0168-1 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: INCYTE PHARMACEUTICALS, INC. 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Word Perfect 6.1/MS-DOS 6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/075,454
FILING DATE: Herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: RAS PROTEINS NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                 Sequence 11, Application US/09075454
Patent No. 6391580
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                           APPLICANT: Patterson, Chandra
APPLICANT: Batra, Sajeev
APPLICANT: Baughn, Mariah R.
TITLE OF INVENTION: RAS PROTE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: 650-855-0555
                                                                                                                                                                                                                                         Tang, Y. Tom
Lal, Preeti
Guegler, Karl J.
Corley, Neil C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.5%;
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
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Search completed: January 16, 2003, 02:47:35 Job time: 87 secs

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January 16, 2003, 02:47:19 ; Search time 283 Seconds (without alignments) 5131.424 Million cell updates/sec
                                                                                                                                                                          GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                    OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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length: 2000000000
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Maximum DB seq
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Perfect score:
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                                                                                                       Run on:
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

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2: /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
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4: /cgn2_6/ptodata/2/pubpna/US00_PUBCOMB.seq:* Published_Applications_NA:* 66... 1111... 14... Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Sequence 1, Appli		Sequence 88, Appl				Sequence 507, App	Sequence 31, Appl		Sequence 303, App	Sequence 26, Appl	Sequence 1930, Ap	Sequence 1930, Ap	Sequence 9, Appli	Sequence 7, Appli	Sequence 879, App	Sequence 885, App	Sequence 894, App	Sequence 896, App
	ID	US-09-817-198A-1	US-09-817-198A-3	US-09-764-868-88	US-09-817-198A-28	US-09-817-198A-29	US-09-817-198A-30	US-09-764-868-507	US-09-817-198A-31	US-09-920-300A-303	US-10-033-528-303	US-09-964-824A-26	US-10-046-935-1930 '	US-09-878-178-1930	US-09-794-257-9	US-09-794-257-7	US-09-834-975-879	US-09-834-975-885	US-09-834-975-894	US-09-834-975-896
		10	10	σ	10	10	10	σ	10	10	12	10	σ	6	10	10	10	10	10	10
	Query Match Length DB	3257	28770	2021	601	601	601	266	601	481	481	463	310	310	624	1161	2497	2497	2497	2497
æ	Query Match	100.0	85.8	35.6	18.4	18.4	16.6	16.4	15.9	14.7	14.7	13.5	9.4	9.4	6.7	6.7	6.7	6.7	6.7	6.7
	Score	3257	2696.6	1160.6	599	599	540	534.2	516.6	477.4	477.4	438.8	306.8	306.8	217.8	217.8	217.8	217.8	217.8	217.8
	Result No.	1	7	3	4	S	9	7	œ	σ	10	c 11	12	13	14	15	16	17	18	19

Sequence 32, Appl Sequence 631, Appl Sequence 816, Appl Sequence 816, Appl Sequence 1359, Appl Sequence 1139, Appl Sequence 141, Appl Sequence 11, Appl Sequence 11, Appl Sequence 14, Appl Sequence 17, Appl Sequence 71, Appl Sequence 71, Appl Sequence 529, Appl Sequence 1703, Appl Sequence 835, Appl Sequence 835, Appl	
US-09-817-198A-32 US-09-925-300-631 US-09-925-300-631 US-09-918-818-836 US-09-918-81-1359 US-09-917-800A-1461 US-09-917-800A-1461 US-09-917-800A-1461 US-09-917-800A-1461 US-09-917-800A-1461 US-09-917-800A-1461 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-764-868-71 US-09-770-445-529 US-09-770-445-529 US-09-917-800A-1426 US-09-917-800A-1426 US-09-918-842A-2113 US-09-938-842A-2113	
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ALIGNMENTS

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        Sequence 1, Application US/09817198A
Patent No. US20020146758A1
GENERAL INFORMATION:
APPLICANT: YE, Jane et al.
TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
FILE REPERENCE: CLO01188
CURRENT APPLICATION NUMBER: US/09/817,198A
CURRENT APPLICATION NUMBER: US/09/817,198A
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                            GGCAGGAGAGATACCAGACCATCACAAAGCAGTACTATCGGCGGGCCCAGGGGATATTTT 300
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100.0%; Pred. No. 0;
Live 0; Mismatches
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Best Local Similarity
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US-09-817-198A-1
US-09-817-198A-1
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Patent No. US20020146758A1

GENERAL INFORMATION:
APPLICAMY: VE. Jane et al.
TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THE.
TITLE OF INVENTION: PROTEINS, AND USES THEREOF
FILE REFERENCE: CLO01188
CURRENT APPLICATION NUMBER: US/09/817,198A
CURRENT APPLICATION NUMBER: 2001-03-27
NUMBER OF SEQ ID NOS: 33
SEQ ID NOS: 33
SEQ ID NOS: 33
SEQ ID NOS: 33
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                                          Length
                                                                       Indels
                                          10;
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                                          Score 2696.6;
Pred. No. 0;
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Matches 2717; Conser
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US-09-817-198A-3
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27602 AGTIGTAAATCACTGGCTAATGAGAAAGGAGACAGCTAACTCTAGGATGAAGCTGTGAC 27661
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0; Mismatches 9;
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TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PTZ32
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CURRENT FILING DATE: 2001-01-17
Prior application data removed - refer to
NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: Patentin Ver. 2.0
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Patent No. US20020168711A1
GENERAL INFORMATION:
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Pred. No. 5.8e-117;
1; Mismatches 1;
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ANISM: Homo sapien
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PT232

FURRENT APPLICATION NUMBER: US/09/764,868

CURRENT FILING DATE: 2001-01-17

Prior application data removed - refer to PALM or file wrapper
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                       GENERAL INFORMATION:
APPLICANT: YE, Jane et al.
TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING
TITLE OF INVENTION: PROTEINS, AND USES THEREOF
FILE REFERENCE: CL001188
CURRENT APPLICATION NUMBER: US/09/817,198A
CURRENT FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16.6%; Score 540; DB 10; 99.6%; Pred. No. 1.5e-104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Mismatches
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Matches 540; Conservative
US20020146758A1
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LENGTH: 601
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 601;
                                                                                   Sequence 29, Application US/09817198A
Patent No. US20020146758A1
GENERAL INFORMATION:
TTLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THES
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THES
TITLE OF INVENTION: OF ORDER 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ISULT 6
5-09-817-198A-30
Sequence 30, Application US/09817198A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ORGANISM: Homo sapien
US-09-817-198A-29
                                                        -09-817-198A-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Simi
Matches 599;
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2001-03-27
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LOCATION: 417, 461
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                                                                 TYPE: DNA
ORGANISM: Homo sapien
 CURRENT FILING DATE:
                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-920-300A-303
                                                                                         US-09-817-198A-31
                                                                                                                                              Matches 516;
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LENGIH: 481
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                                                                                                                     Query Match
Best Local
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Patent No. US20020146758A1

GENERAL INFORMATION:

APPLICANT: YE, Jane et al.

TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,

TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE

TITLE OF INVENTION: PROTEINS, AND USES THEREOF

FILE REFERENCE: CL001188

CURRENT APPLICATION NUMBER: US/09/817,198A
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; OTHER INFORMATION: n equals a,t,g,
US-09-764-868-507
                                                                                                                     equals a,t,g,
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                                                                                                                                                                                                  equals
NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                           Best Local Similarity 98.9
Matches 533; Conservative
                                   LENGTH: 566
TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: SITE
LOCATION: (484)
OTHER INFORMATION: n equi
                                                                                                                                            LOCATION: (493)
OTHER INFORMATION: n
NAME/KEY: SITE
LOCATION: (538)
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                        ID NO 507
ENGTH: 566
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            SOFTWARE:
SEQ ID NO 5
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TCCAGCCCCAGGCACGCCACCTGCTCCTGAGCCTCCACTATCTCCCTGTGACGGGTGAAC 3126
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APPLICANT: Meagher, Madeleine Joy
APPLICANT: Xu, Jianghoun
APPLICANT: Xu, Jianghoun
APPLICANT: Xu, Jianghoun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TITLE OF INVENTION: COMPOSITIONS OF COLON CANCER
FILE REPERENCE: 210121:547
CURRENT APPLICATION NUMBER: US/09/920,300A
CURRENT FILING DATE: 2001-07-31
NUMBER OF SEQ ID NOS: 1789
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                DB 10;
                                                                                                                                                                                                                Score 516.6; DB 10;
Pred. No. 1.2e-99;
; Mismatches 0;
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NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 31
LENGTH: 601
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ilarity 99.8%;
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GENERAL INFORMATION:
APPLICANT: HOTRIGAM. Stephen
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using Sign
TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using AFLIS OF TOWN UNMER: US/60/2964,824A
CURRENT PILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
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421 GITAACCAGGCTCTGCTGGTGTTACTCGTCACCACTNIGCACCTGCTGTTTAGAGA 480
                                                                                                             Gaps
                                                                                                                                                                    GTTGCTTCCTTGAAGATGGGACTCCTTGGGTATCAAGACCTATGCCACATCACACTGGGG
                                                                                             CTAGGGAAGTAGGTGATGCCAGCCCTCAAGTCTGTCTTCAGCCAGGGACTTGAGAAGTTA
                                                                                                                                                                                                                                                                     GTACCCTAGGGGTTCTGGAAGGGAAACATGGAATCAGGATTCTATAGACTGATAGGCCC
                                                                                                                                                                                                                                                                                                                                                241 GTACCCTAGGGGTTCTGGAAGGAAAACATGGAATCAGGATCTATAGACTGATAGGCCC
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                                  GGTGCCATTCATTGTCTTTCTCTCCTAGCCCCCTCAGGAAAGAAGGACTATTTTGTACT
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Pred. No. 2.5e-83;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 26, Application US/09964824A Patent No. US20020102531A1
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ilarity 99.3%;
Conservative (
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US-09-964-824A-26/c
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LENGTH: 463
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APPLICANT: King, Gordon E.
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Meagher, Madeleine Joy
APPLICANT: Au, Jiangchun
APPLICANT: Secrist, Heather
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER
FILE REPERENCE: J10121.54701
CURRENT APPLICATION NUMBER: US/10/033,528
CURRENT FILING DATE: 2001-12-26
SOFTWARE: FastSEQ for Windows Version 4.0
                                                          Length
                                                                                               Indels
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99.4%; Pred. No. 2e-91;
Live 0; Mismatches 3; ]
                                                        Score 477.4; DB 10;
Pred. No. 2e-91;
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CCATION: 417, 461

COTHER INFORMATION: n = A,T,C or

US-10-033-528-303
    or
                                                      Query Match
Best Local Similarity 99.4%;
Matches 478; Conservative
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Matches 478; Conservative
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; OTHER INFORMATION: n
US-09-920-300A-303
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; ORGANISM: Homo sapien
US-09-878-178-1930
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US-09-878-178-1930
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Fatent No. US20020156011A1

GENERAL INFORMATION:
APPLICANT: Harlocker, Susan L.
APPLICANT: Secrist, Heather
APPLICANT: Secrist, Heather
APPLICANT: Stolk, John A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE T TITLE OF INVENTION: COMPOSITIONS OF COLON CANCER
FILE REFERENCE: 210121.527C1
CURRENT FILIG DATE: 2002-01-15
CURRENT FILIG DATE: 2002-01-15
NUMBER OF SEQ ID NOS: 2239
SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 310
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Matches 308; Conservative
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; ORGANISM: Homo sapiens
US-10-046-935-1930
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Sequence 9, Application US/09794257

Patent No. US20020009804A1

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: 32705, 23224, 27423, 32700, 32712, No. US20020009804A1e1

TITLE OF INVENTION: Human G-Proteins
FILE REFERENCE: 35800/203285

CURRENT APPLICATION NUMBER: US/09/794,257

CURRENT FILING DATE: 2001-02-27

PRIOR FILING DATE: 2000-02-29
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Sequence 1930, Application US/09878178

Fatent No. 052002017752A1

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Jiang, Yugiu

APPLICANT: Secrist, Heather

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

TITLE OF INVENTION: AND DIAGNOSIS OF COLON CANCER

FILE REFERENCE: 210.11.527

CURRENT APPLICATION NUMBER: US/09/878,178

CURRENT APPLICATION NUMBER: US/09/878,178

CURRENT FILING DATE: 2001-06-08

NUMBER OF SEQ ID NOS: 2237

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NOS: 2237
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SOFTWARE: FastSEQ for Windows Version 4.0
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Sequence 7, Application US/09794257

Patent No. US20020009804a1

GENERAL INFORMATION:
APPLICANT: Weapers, Rachel
TITLE OF INVENTION: 32705, 23224, 27423, 32700, 32712, No. US20020009804a1e1
TITLE OF INVENTION: Human G-Proteins
FILE REFERENCE: 35800/220985

CURRENT APPLICATION NUMBER: 05/09/794,257

CURRENT FILING DATE: 2001-02-27

PRIOR APPLICATION NUMBER: 60/185,606

PRIOR FILING DATE: 2000-02-29

NUMBER OF SEQ ID NOS: 16

SCOFTWARE: FastSEQ for Windows Version 4.0
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                                                  Length 624;
                                                Score 217.8; DB 10; Length
Pred. No. 8e-37;
0; Mismatches 167; Indels
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Pred. No. 9.9e-37;
0; Mismatches 167;
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ilarity 65.6%;
Conservative C
                                                 6.78;
                                              Query Match
Best Local Similarity 65.65
Matches 318; Conservative
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; LOCATION: (18)...(641)
US-09-794-257-7
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ORGANISM: homo sapiens
TYPE: DNA ORGANISM: homo sapiens
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Matches 318; Conserv
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                      US-09-794-257-9
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US-09-794-257-7
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           18 ATGGCGAAGACGTACGATTATCTCTTCAAGCTCCTGCTGATCGGCGACTCGGGGGTAGGC
                                         AAGACCTGCCTGCTGTGCCGCTTCACCGACAACGAGTTCCACCTCCTCGCACATCTCCACC
                                                                                                                              ATCTGGGACACTGCAGGGCAGGAGAGATACCAGACCATCACAAAGCAGTACTATCGGCGG
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                                                                                   ATCGGTGTTGACTTTAAGATGAAGACCATAGAGGTAGACGGCATCAAAGTGCGGATACAG
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Perfect score: Sequence: Scoring table:

Run on:

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Database

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Sequence E Sequence 1 Sequence 1

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49.3%; Score 545; DB 2; Length 207; ilarity 52.9%; Pred. No. 2.2e-52; Conservative 45; Mismatches 43; Indels
                                Sequence 4, Application US/08824873
; Sequence 4, Application US/08824873
; Patent No. 5843710;
; GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Geogler, Karl
TITLE OF INVENTION: NOVEL RAB PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS;
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STRRET: 3174 Porter Drive
CITY: Palo Alto
STAPE: CA
COUNTRY: USA
ZIP: C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-718-270A-11
US-08-531-525-26
US-08-718-270A-26
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SOFTWARE: FastSED for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/824,873
FILING DATE: Filed Herewith
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NAME: B1111ngs, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/COCKET NUMBER: PF-0:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-85-0555
TELEPHONE: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
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Best Local Similarity
Matches 99; Conserv
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LIBRARY: Generated Control Cont
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY:
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TYPE: am
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                                                                                                                                                                                                                            January 16, 2003, 02:47:39; Search time 24 Seconds (without alignments) 259.902 Million cell updates/sec
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Sequence 3
Sequence 6
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1 MAKQYDVLFRLLLIGDSGVG......LEBEEGKPEGPANSSKTCWC
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/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
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Copyright (c) 1993 - 2003 Compugen Ltd.
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US-08-718-270A-51
US-08-511-525-51
US-08-718-270A-35
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US-08-718-270A-10
US-08-718-270A-14
US-08-718-270A-34
US-09-116-901-3
US-09-116-901-8
US-09-116-901-8
US-09-116-901-8
US-08-118-270A-14
US-08-511-525-13
US-08-718-270A-13
US-08-718-270A-13
US-08-718-270A-13
US-08-718-270A-13
US-08-198-184-3
US-08-198-184-3
US-08-198-184-3
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US-08-824-873-1
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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453.5 453.5 452.5 452.5 443

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123 ADEEOKROVGREGGOQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGLRM 182
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61 IWDTAGQERFRTITTAYYRGAMGIMLVYDITNEKSFDNIRNWIRNIEEHASADVEKMILG 120
                                    121 NKADEEQKROVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGL 180
                                                       47.7%; Score 527.5; DB 2; Length 205; 52.4%; Pred. No. 1.9e-50;
                                                                                                                                                                                                                                                                                                                                                                             Peptides Inhibiting the Oncogenic Action of P21 Ras
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SULTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DAPA: APPLICATION NUMBER: US/OB/531 6.7 FILLE FILLED NAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.9e-50;
hes 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Greenlee and Winner, P.C.
5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                        APPLICANT: Pincus, Matthew R.
APPLICANT: No. 58406831e, John F.
APPLICANT: Abajian, Henry B.
APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Peptides Inhibiti TITLE OF INVENTION: Of P21 Ras
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSE: Greenics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 21-SEP-1995
CLASSIFICATION: 530
TORNEY AACTORN
                                                                                                                                                                                                                                         Sequence 25, Application US/08531525
Patent No. 5840683
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: FETEA. Donna M.
REGISTRATION NUMBER: 33.87-9
TELECOMUNICATION INFORMATION:
TELEPHONE: (303) 499-8089
INFORMATION FOR SEQ ID NO: 25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            45;
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
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Best Local Similarity 52.43
Matches 97; Conservative
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MOLECULE TYPE: protein
HYPOTHETICAL: NO
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ORGANISM: Can
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                                                                                                                                                       NKADEEOKROVGREOGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGL 180
                                                                                                                                                                            61 IWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIG 120
                                                                                                   61 IWDTAGGERFRITTAYYRGAMGIMLVYDITUEKSFDNIRNWIRNIEEHASADVEKMILG 120
                              1 MAKQYDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    49.3%; Score 545; DB 3;
ilarity 52.9%; Pred. No. 2.2e-52;
Conservative 45; Mismatches 43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSED for Windows Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/198,184
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Hillman, Jennifer L.
APPLICANT: Guegler, Karl
TITLE OF INVENTION: NOVEL RAB PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STRRET: 3174 Porter Drive
CITY: Palo Alto
STATE: CA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PF-0240 US
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APPLICATION NUMBER: 08/824,873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTATION UNDRER: 36,749
REFERENCE/DOCKET NUMBER: PF-02
REFERENCE/DOCKET NUMBER: PF-02
REFERENCE/TOOKET NUMBER: PF-02
REFERENCE/TOOKET NUMBER: PF-02
REFERENCE/TOOKET NUMBER: PF-02
RELEFAX: 415-845-055
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                              Sequence 4, Application US/09198184 Patent No. 6010859 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 207 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss: single
linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
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CLONE: 234746
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Matches 99; Conserv
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IMMEDIATE SOURCE:
                                                                                                                                                                                                                                     181 RMRASNE 187
                                                                                                                                                                                                                                                                          181 SPQGSNQ 187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
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123 ADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGLRM 182
                         61 IWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIG 120
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                                                                                                                                                                                                                Sequence 51, Application US/08531525
Patent No. 5840683
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Pincus, Matthew R.
APPLICANT: No. 58406811e, John F.
APPLICANT: Abajian, Henry B.
APPLICANT: Abajian, Henry S.
TITLE OF INVENTION: Peptides Inhibiting the Oncogenic Action TITLE OF INVENTION: of P21 Ras
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 198;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121 NKADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQ 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: LLM. COMPUTER: COMPUTER: COMPUTER: COPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/531,525 FILING DATE: 21-SEP-1995 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     46.9%; Score 518.5; DB 2
55.0%; Pred. No. 1.8e-49;
Live 45; Mismatches 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3: Greenlee and Winner, P.C.
5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY AGENT INFORMATION:
NAME: Ferber, Donna M.
REGISTATION NUMBER: 37.878
REFERENCE/DOCKET NUMBER: 37-
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: (303) 499-8080
TELEFAX: (303) 499-8089
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 55.0%
Matches 94; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein HYPOTHETICAL: NO
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) ORGANISM: Can

US-08-531-525-51
                                                                                                                        181 QGSNQ 185
                                                                                  183 RASNE 187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 KQYDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQIW 62
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                                                                                                                                                                                                                                        APPLICANT: Hlavka, Joseph J.
APPLICANT: Pincus, Matthew R.
APPLICANT: No. 5910478le, John F.
APPLICANT: Abajian, Henry B.
APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Peptidomimetics Inhibiting
TITLE OF INVENTION: the Oncogenic Action of P21 Ras
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             42;
                                                                                                                                                                                                                                                                                                                                                                                                                               Greenlee, Winner and Sullivan, P.C.
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52.4%; Pred. No. 1.9e-50;
Live 45; Mismatches 42
                                                                                                                                                                                                                                                                                                                                                                                                                                              5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DAYR:
APPLICATION NUMBER: US/08/718,270A
FILING DATE: 20-SEP-1996
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/531,525
FILING DATE: 21-SEP-1995
FILING DATE: 21-SEP-1995
FILING DATE: 21-SEP-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,878
RÉFERENCE/DOCKET NUMBER: 78-95
TELECOMMUNICATION INFORMATION:
TELEPHONE: (303) 499-8080
TELEPAX: (303) 499-8089
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                             Sequence 25, Application US/08718270A Patent No. 5910478 GENERAL INFORMATION:
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amino acid
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Matches 97; Conservative
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CORRESPONDENCE ADDRESS:
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MOLECULE TYPE: pr
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                                                                 : ||:
181 QGSNQ 185
                                        183 RASNE 187
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US-08-718-270A-25
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                                                                                                                                                          US-08-718-270A-25
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183 RASNELALAE-LEEEEGKPEGPANSSK 208
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             ; Sequence 35, Application US/08531525
; Patent No. 5840683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY AGENT INFORMATION:
NAME: Ferber, Donna M.
REGISTATION UNDRER: 33,878
REFERENCE/DOCKET UNBER: 37-94
TELECOMMUNICATION INFORMATION:
TELEPAN: (303) 499-8080
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
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                                              GENERAL INFORMATION:
APPLICANT: Hlavka, Joseph J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 207 amino acids
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                                                                                                                                                                                               NUMBER OF SEQUENCES:
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Matches 96; Conserv
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STRANDEDNESS:
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ORGANISM: Dis
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US-08-531-525-35
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                                                                                                                                                                                  Peptidomimetics Inhibiting
the Oncogenic Action of P21 Ras
52
                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/718,270A FILING DATE: 20-SEP-1996 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
                                                                                                                                                                                                                                                           ADDRESSEE: Greenlee, Winner and Sullivan, P.C. STREET: 5370 Manhattan Circle, Suite 201 CITY: Boulder STATE: Colorado
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APPLICATION NUMBER: US 08/531,525

FILING DATE: 21-SEP-1995

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 60/004,091

FILING DATE: 21-SEP-1995

ATTORNEY/AGENT INFORMATION:
                                                                                                       APPLICANT: Hlavka, Joseph J.
APPLICANT: Pincus, Matthew R.
APPLICANT: No. 59104781e, John F.
APPLICANT: Abajian, Henry B.
APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Peptidomimetics
TITLE OF INVENTION: the Oncogenic A
                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                    Sequence 51, Application US/08718270A Patent No. 5910478 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         78-95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Ferber Donna M.
REGISTATION NUMBER: 33,878
REFERENCE/DOCKET NUMBER: 78-95
REFCOMMUNICATION INFORMATION:
TELEPHONE: (303) 499-8080
TELEFAX: (303) 499-8080
INFORMATION FOR SEQ ID NO: 51:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Canis familiaris
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Matches 94; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 198 amino acida
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MOLECULE TYPE: protein
HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
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STRANDEDNESS: si
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                                                                                                                                                                                                                                                                                                                                                             80303
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                      RESULT 6
US-08-718-270A-51
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63 DTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIGNK 122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           123 ADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGLRM 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3 KQYDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQIW 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 KTYDYLFKLLLIGDSGVGKTCLLFRFSEDAFNTTFISTIGIDFKIRTVELDGKKIKLQIM 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11;
APPLICANT: Pincus, Matthew R.
APPLICANT: No. 58406831e, John F.
APPLICANT: Abajian, Henry B.
APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Peptides Inhibiting the Oncogenic Action TITLE OF INVENTION: of P21 Ras
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 207;
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                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION DATA:
APPLICATION NUMBER: US/08/531,525
FILING DATE: 21-SEP-1995
CLASSIFICATION: S30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              45.6%; Score 503.5; DB 2 ilarity 46.4%; Pred. No. 8.8e-48; Conservative 56; Mismatches 44
                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Greenlee and Winner, P.C.
STREET: 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 8
US-08-718-270A-35
US-08-718-270A-35
Sequence 35, Application US/08718270A
Patent No. 5910478
GENERAL INFORMATION:
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RESULT 7

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65 AGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIGNKAD 124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 YDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQIWDT 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 YDYLIKLLIGDSGVGKSCLLRFSDGSFTTSFITTIGIDFKIRTIELDGKRIKLQIWDT 71
                                                                                                APPLICANT: Pincus, Matthew R.
APPLICANT: No. 58406831e, John F.
APPLICANT: Abajian, Henry B.
APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Peptides Inhibiting the Oncogenic Action NUMBER OF SEQUENCES: 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2; Length 215;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            44.8%; Score 494.5; DB 2
illarity 46.9%; Pred. No. 9.2e-47;
Conservative 40; Mismatches 54
                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Greenlee and Winner, P.C.
STREET: 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             180 LSDIDSRAEPATIKISQIDQAAGAGQATQKSAC 212
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/531,525
FILING DATE: 21-SEP-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Ferber, Donna M.
REGISTATION NUMBER: 37-94
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: (303) 499-8089
INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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Sequence 10, Application US/08718270A
Patent No. 5910478
GENERL INFORMATION:
APPLICANT: Hlavka, Joseph J.
APPLICANT: Pincus, Matthew R.
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                                                                         Hlavka, Joseph J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 215 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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Best Local Similarity
Matches 100; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Boulder
STATE: Colorado
                                         INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORIGINAL SOURCE
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                                                                   APPLICANT:
APPLICANT:
APPLICANT:
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                                         GENERAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2; Length 207;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE PATENTIN STATES OF THE SOFTWARE SOFTWAR
                                                                                                                         APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Peptidomimetics Inhibiting
TITLE OF INVENTION: the Oncogenic Action of P21 Ras
NUMBER OF SEQUENCES: 52
                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Greenlee, Winner and Sullivan, P.C. STREET: 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             45.6%; Score 503.5; DB 2
46.4%; Pred. No. 8.8e-48;
Live 56; Mismatches 44
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
Hlavka, Joseph J.
Pincus, Matthew R.
No. 59104781e, John F.
Abajian, Henry B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Ferber, Donna M.
REGISTRATION NUMBER: 33,878
REFERENCE/DOCKET NUMBER: 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (303) 499-8089
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 207 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                Boulder
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                                                                                                                                                                                                                                                                                                                                                                                                                                    80303
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                                                                                                                                                                                                                                                                                                                                                                    STATE: C.
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US-08-531-525-10
                                                                                         APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Matches 9
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------RDIKQR 179

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60 QIWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILI 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Peptides Inhibiting the Oncogenic Action
TITLE OF INVENTION: of P21 Ras
NUMBER OF SEQUENCES: 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 194;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             120 GNKADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQ 171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Pincus, Matthew R.
APPLICANT: No. 59104781e, John F.
APPLICANT: Abajian, Henry B.
APPLICANT: Rende, Andrew S.
APPLICANT: Kende, Andrew S.
APPLICANT: Kende, Andrew S.
APPLICANT: Kende, Andrew S.
A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/531,525 FILING DATE: 21-SEP-1995 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    44.2%; Score 488.5; DB 2; 55.8%; Pred. No. 3.6e-46;
                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Greenlee and Winner, P.C.
STREET: 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 41; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 80303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 34, Application US/08718270A Patent No. 5910478
GENERAL INFORMATION:
APPLICANT: Hlavka, Joseph J. APPLICANT: Pincus, Matthew R. APPLICANT: No. 59104781e, John F.
Hlavka, Joseph J.
Pincus, Matthew R.
No. 58406831e, John F.
Abajian, Henry B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Ferber, Donna M.
REGISTRATION UNDRER: 33,878
REFERENCE/DOCKET UNDRER: 37-9.
TELECOMMUNICATION INFORMATION:
TELEPHOR: (303) 499-8080
TELEFAX: (303) 499-8089
INFORMATION FOR SEQ ID NO: 34:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORIGINAL SOURCE:
ORGANISM: Discopyge ommata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 194 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 55.88
Matches 96; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                           Colorado
: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 21 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                      Boulder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
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US-08-718-270A-34
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                                                                                           APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                    APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  qq
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIGNKAD 124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12 YDYLIKLELIGDSGVGKSCLLRFSDGSFTTSFITTIGIDFKIRTIELDGKRIKLQIWDT 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 YDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQIWDT 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 215;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/718,270A
FILING DATE: 20-SEP-1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/531,525
FILING DATE: 21-SEP-1995
PRIOR APPLICATION NUMBER: US 60/004,091
APPLICATION NUMBER: US 60/004,091
FILING DATE: 21-SEP-1995
ATTOMEX/AGENT IRFORMATION:
                                    APPLICANT: Abajian, Henry B.
APPLICANT: Kende, Andrew S.
TITLE OF INVENTION: Pettidomimetics Inhibiting
TITLE OF INVENTION: the Oncogenic Action of P21 Ras
NUMBER OF SEQUENCES: 52
CORRESPONDENCE ADDRESS:
ADDRESSE: Greenlee, Winner and Sullivan, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 2;
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                                                                                                                                                                                                            E: Greenlee, Winner and Sullivan, P.C. 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           44.8%; Score 494.5; DB
46.9%; Pred. No. 9.2e-47
Live 40; Mismatches 5
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 34, Application US/08531525
Patent No. 5840683
GENERAL INFORMATION:
      No. 59104781e, John F. Habajian, Henry B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Arabidopsis thaliana
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Ferber, Donna M. REGISTRATION NUMBER: 33,878 REFERENCE/DOCKET NUMBER: 78-TELECOMMUNICATION INFORMATION: (303) 499-8080
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (333) 499-8089 INFORMATION FOR SEQ ID NO: 1 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 46.9%;
Matches 100; Conservative
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MOLECULE TYPE: protein
HYPOTHETICAL: NO
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CITY: Boulder
STATE: Colorado
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORIGINAL SOURCE:
ORGANISM: Ara
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TELEFAX: (
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US-08-531-525-34
                                                                                                                                                                                                                                                                                                                                   COUNTRY:
            APPLICANT:
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61 IWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        121 NKADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              121 NKSDLTTKKVVDNTTAKEFADSLGIPFLETSAKNATNVEQAF-----MTMAAEIKKRMGP 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 MAKQYDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQ 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Score 478.5; DB 2; Pred. No. 4.9e-45; 42; Mismatches 64
                                                                                                                          SYSTEM: DOS
FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: Incyte Pharmaceuticals, Inc 3174 Porter Dr.
                                                                                                                                                                                                                                                                                                                                                      PF-0367 US
                                                                                                                                                                             APPLICATION NUMBER: US/08/916,901 FILING DATE: Filed Herewith PRIOR APPLICATION DATA: APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 3, Application US/09154602
Patent No. 6300472
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Carley, Neil C.
APPLICANT: Carley, Neil C.
APPLICANT: Shah, Purvi
TITLE OF INVENTION: RAB PROTEINS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: BILLINGS, LUCY J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-85-0555
TELEPHONE: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      181 RMRASNELALAELEEEGKPEG 202
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                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 45.0%;
Matches 91; Conservative 4;
                                                                                                   COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          201 amino acids
                                                                                                                                                              CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; LIBRARY: LIVRTUTO4
; CLONE: 2514506
US-08-916-901-3
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                                                                                                                          OPERATING SYSTEM:
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                       USA
                       COUNTRY: US
ZIP: 94304
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US-09-154-602-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 194;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     120 GNKADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQ 171
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                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 44.2%; Score 488.5; DB 2; Best Local Similarity 55.8%; Pred. No. 3.6e-46; Matches 96; Conservative 41; Mismatches 30;
                   5: Greenlee, Winner and Sullivan, P.C. 5370 Manhattan Circle, Suite 201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Corley, Neil C.
APPLICANT: Shah, Purvi
ITTLE OF INVENTION: RAB PROTEINS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/718,270A FILING DATE: 20-SEP-1996 CLASSPICATION: 514 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/531,525 PRIOR APPLICATION DATA: APPLICATION NUMBER: US 60/004,091 FILING DATE: 21-SEP-1995 ATTORNEY/AGENT INFORMATION:
                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33,878
33,878
--- 78-95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Ferber, Donna M.
REGIETRATION UNDHBER: 33,878
RECERRATION UNDHBER: 78-9
TELECOMMUNICATION INFORMATION:
TELEPAN: (303) 499-8089
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-916-901-3; Sequence 3, Application US/08916901; Patent No. 5892012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US:08-718-270A-34 Discopyge ommata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  194 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                         CITY: Boulder
STATE: Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Palo Alto
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ORGANISM: Dis
                                                                                                                  80303
                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dp
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Length 201; Indels

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Search completed: January 16, 2003, 04:18:07 Job time : 26 secs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  121 NKADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MAKQYDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQ 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 201;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
43.3%; Score 478.5; DB 4
Best Local Similarity 45.0%; Pred. No. 4.9e-45;
Matches 91; Conservative 42; Mismatches 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FSSLSED for Windows Version 2.0
CURRENT APPLICATION DARB: US/08/916,901
                    SYSTEM: DOS
FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Incyte Pharmaceuticals, Inc
                                                                                                                                                                                                                  PF-0367 US
                                     SOFTWARE: FASTSEQ for Windows Ver:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/154,602
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hillman, Jennifer L.
APPLICANT: Lal, Preeti
APPLICANT: Corley, Neil C.
APPLICANT: Shah, Purvi
TITLE OF INVENTION: RAB PROTEINS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/916,901
FILING DATE:
                                                                                                                                                        ATTOREY AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-03
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-85-055
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 201 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8, Application US/08916901
Patent No. 5892012
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         181 RMRASNELALAELEEEGKPEG 202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          176 GAASGGERPNLKIDSTPVKPAG 197
COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Incyte Pharr
STREET: 3174 Porter Dr
CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                              single
                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: LIVRTUTO4
CLONE: 2514506
                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: sil
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US-08-916-901-8
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61 IWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Score 469; DB 2;
; Pred. No. 5.5e-44;
36; Mismatches 43;
                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0367 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEPHONE: 415-845-4166
INFORMATION FOR SEQ ID NO: 8:
FILING DATE: Filed Herewith PRIOR APPLICATION DATA: APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    42.4%;
ilarity 51.2%;
Conservative 3
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LENGTH: 201 amino acids
TYPE: amino acid
STRANDEDNESS: single
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linear
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LIBRARY: GenBank
CLONE: 57006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
nes 83; Conserv
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January 16, 2003, 04:18:15; Search time 77 Seconds (without alignments) 54.730 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                             1105
1 MAKQYDVLFRLLLIGDSGVG......LEEEEGKPEGPANSSKTCWC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1: /cgn2_6/ptodata/2/pubpaa/USO8_NEW_PUB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
4: /cgn2_6/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*
5: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
6: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
7: /cgn2_6/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
8: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*
9: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*
10: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*
11: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*
11: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*
13: /cgn2_6/ptodata/2/pubpaa/USO9_NEW_PUB.pep:*
13: /cgn2_6/ptodata/2/pubpaa/USO0_NEW_PUB.pep:*
13: /cgn2_6/ptodata/2/pubpaa/USO0_NEW_PUB.pep:*
14: /cgn2_6/ptodata/2/pubpaa/USO0_NEW_PUB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           120991 segs, 19878514 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Published_Applications_AA:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                   OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                    BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 length: 0
length: 2000000000
                                                                                                                                                                                                                                                                                                 US-09-817-198A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Minimum DB seq
Maximum DB seq
                                                                                                                                                                                                                                                                                              Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
                                                                                                                                                                                                                                                                                                                                                         Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Database
                                                                                                                                                                           Run on:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	Sequence 2, Appli	Sequence 701, App	Sequence 5, Appli		Sequence 1120, Ap	Seguence 534, App	Sequence 1571, Ap	Sequence 8, Appli	Sequence 5, Appli	Sequence 766, App	Sequence 3, Appli	æ		Sequence 16, Appl	Sequence 5, Appli	4	Sequence 4, Appli	7	84,
ID	US-09-817-198A-2	US-09-764-868-701	US-09-817-198A-5	US-09-817-198A-4	US-09-764-868-1120	US-09-925-302-534	US-09-925-300-1571	US-09-794-257-8	US-09-822-860-5	US-09-834-765-766	US-09-967-736-3	US-09-967-736-8	US-09-820-003A-4	US-09-794-257-16	US-09-945-173-5	US-09-972-529-4	US-09-817-199A-4	US-09-817-199A-2	US-09-764-868-684
DB	10	6	10	10	6	10	10	10	10	10	10	10	10	10	10	10	10	10	δ
% Query Match Length DB	212	401	218	212	188	246	218	207	190	162	201	201	222	198	198	198	223	223	226
% Query Match	100.0	100.0	98.8	97.5	75.3	48.1	48.0	47.8	46.3	45.0	43.3	42.4	42.3	39.3	39.3	39.3	38.9	38.8	38.6
Score	1105	1105	1092	1077	832	532	530.5	528.5	512	497	478.5	469	467	434.5	434.5	434.5	430	429	427
Result No.	1	7	m	4	Ŋ	9	7	80	6	10	11	12	13	14	15	16	17	18	19

RESULT 2

Sequence 1106, Ap Sequence 14, Appl Sequence 3, Appli Sequence 67, Appl	Sequence 1100, Ap Sequence 783, App Sequence 5, Appli Sequence 1112, Ap Sequence 692, App	Sequence 8, Appli Sequence 3, Appli Sequence 1077, Ap Sequence 2, Appli	Sequence 5, Appli Sequence 762, App Sequence 2, Appli Sequence 765, App Sequence 763, App Sequence 763, App	Sequence 10, Appl Sequence 1364, Ap Sequence 6, Appli Sequence 2, Appli Sequence 1, Appli Sequence 4, Appli Sequence 700, App
9 US-09-764-868-1106 10 US-09-794-257-14 12 US-10-051-986-3 10 US-09-350-874-67	9 US-09-764 868-1100 10 US-09-925-302-783 10 US-09-794-257-5 9 US-09-764-868-1112 9 US-09-764-868-692	10 US-09-988-974-8 10 US-09-988-974-3 10 US-09-925-301-1077 10 US-09-822-860-2	10 US-09-834-765-5 10 US-09-834-765-762 10 US-09-834-765-2 10 US-09-834-765-765 10 US-09-834-765-765 9 US-10-108-605-45	10 US-09-945-173-10 10 US-09-925-300-1364 12 US-10-051-986-6 10 US-09-820-003A-2 12 US-10-051-986-1 12 US-10-051-986-4 9 US-09-764-868-700
222 191 191 212	307 312 213 222 225	213 217 239 201	624 625 832 168 208	216 217 211 173 259 260 287
38.6 37.6 37.6 37.3	36.1 36.1 34.0 33.9	8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	32.0 32.0 31.8 31.0	30.6 29.0 29.0 29.0 29.0
426 416 416 412.5	399 399 376 374.5	373 373 373	353.5 353.5 353.5 351 343 339	338.5 338.5 323.5 321 320 320 320
20 21 22 23	2002 2002 2002 2008	333	33 34 35 37 38	39 44 44 44 45

ALIGNMENTS

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                                              GENERAL INFORMATION:
APPLICANT: YE, Jane et al.
TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
TITLE OF INVENTION: PROTEINS, AND USES THEREOF
FILE REFERENCE: CL001188
CURRENT FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 IWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                 1 MAKQYDVLFRLLLIGDSGVGKTČLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQ 60
                                                                                                                                                                                                                                                                                                                                                                             ;
0
                                                                                                                                                                                                                                                                                                                                        Query Match 100.0%; Score 1105; DB 10; Length 212; Best Local Similarity 100.0%; Pred. No. 1e-97; Matches 212; Conservative 0; Mismatches 0; Indels 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RMRASNELALAELEEEGKPEGPANSSKTCWC 212
              Sequence 2, Application US/09817198A Patent No. US20020146758A1
                                                                                                                                                                                                                                                                 ; TYPE: PRT
; ORGANISM: Human
US-09-817-198A-2
US-09-817-198A-2
                                                                                                                                                                                                                               SEQ ID NO 2
LENGTH: 212
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Sequence 5, Application US/09817198A

Sequence 5, Application US/09817198A

Patent No. US2002014658A1

GENERAL INFORMATION:
TITLE OF INVENTION: INCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE

TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE

TITLE OF INVENTION: PROTEINS, AND USES THEREOF

FILE REFERENCE: CL001188

CURRENT PELLON NUMBER: US/09/817,198A

CURRENT FILING DATE: 2001-03-27

NUMBER OF SEQ ID NOS: 33

SOCTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                        Length 401;
                                    GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PT232
CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT FILING DATE: 2001-01-17
Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 701
LENGTH: 401
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                                                                                                                                                                                                                                                                                                      Score 1105; DB 9;
Pred. No. 2.3e-97;
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Pred. No. 1.8e-96;
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               Sequence 701, Application US/09764868
Patent No. US20020168711A1
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100.0%;
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Matches 212; Conservative
                                                                                                                                                                                                                                                   ORGANISM: Homo sapiens
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Best Local Similarity
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US-09-764-868-701
                                                                                                                                                                                                                                                                     US-09-764-868-701
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LOCATION: (151)
TOTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids NAME/KEY: SITE
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                                                                                                                                                                                                                                                                               APPLICANT: YE, Jame et al.
TITLE OF INVENTION: ISOLAMED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: MUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
TITLE OF INVENTION: PROTEINS, AND USES THEREOF
FILE REFERENCE: CL001188
CURRENT APPLICATION NUMBER: US/09/817,198A
CURRENT APPLICATION NUMBER: US/09/817,198A
NUMBER OF SEQ ID NOS: 33
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121 NKADEEQKRQVGREQGQQKCPSLQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHR 180
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PT232
CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT APPLICATION GATE: 2001-01-17
Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1120
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                                            Score 1077; DB 10
Pred. No. 4.5e-95;
2; Mismatches 3
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
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97.6%;
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; ORGANISM: Rattus norvegicus
US-09-817-198A-4
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Best Local Similarity
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                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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LOCATION: (139)
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TYPE: PRT
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OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids NAME/KEY: SITE LOCATION: (164)
OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
                                                                            NAME/KEY: SITE

LOCATION: (188)

COTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids US-09-764-868-1120
                                                                                                                                                                                                                                                                                                                                                            61 IWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIG 120
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                                                                                                                                                                                                                                                                                 24 MAKQYDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQ 83
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 514, Application US/09925302
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT APPLICATION NUMBER: US/09/925,302
CURRENT FILING DATE: 2001-08-10
PRIOR PFLICATION NUMBER: PCT/US00/05918
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 1909-03-12
NUMBER OF SEQ ID NOS: 896
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               48.1%; Score 532; DB 10; Length 246; 47.6%; Pred. No. 2.9e-43; tive 49; Mismatches 43; Indels 10
                                                                                                                                                                            Length 188;
                                                                                                                                                                                                                      4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   144 NKADEEQXRQVGREQGQXLAXEYGMDFYETSACTNLNIKESFTR 187
                                                                                                                                                                                                                                                                                                                                                                                                                 121 NKADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTR 164
                                                                                                                                                                              Query Match
75.3%; Score 832; DB 9;
Best Local Similarity 97.6%; Pred. No. 6.9e-72;
Matches 160; Conservative 0; Mismatches 4
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Matches 100; Conservative 4
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ORGANISM: Homo sapiens
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APPLICANT: Meyers, Rachel
TITLE OF INVENTION: 32705, 2324, 27423, 32700, 32712, No. US20020009804Alel
TITLE OF INVENTION: Human G-Proteins
TITLE OF INVENTION: Human G-Proteins
FILE REFERENCE: 35800/209285
CURRENT APPLICATION NUMBER: US/09/794,257
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: 60/185,606
PRIOR FILING DATE: 2000-02-29
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION STATES AND ADDRESS AND ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 48.0%; Score 530.5; DB 10; Best Local Similarity 55.8%; Pred. No. 3.5e-43; Matches 96; Conservative 46; Mismatches 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            33;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8, Application US/09794257 Patent No. US20020009804A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ) ORGANISM: Homo sapiens
US-09-925-300-1571
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: homo sapiens US-09-794-257-8
                                          APPLICANT: Craig Rosen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 95; Conserv
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 1571
LENGTH: 218
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                                                                                                              APPLICANT: ZHU, Shiaoping et al.

TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
TITLE OF INVENTION: PROTEINS, AND USES THEREOF
FILE REFERENCE: CL001214
CURRENT APPLICATION NUMBER: US/09/822,860
CURRENT FILING DATE: 2001-04-02
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         66 GQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIGNKADE 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      126 EQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGLRMRAS 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 DYLFKLLLIGDSGVGKTCLLFRFSEDAFNTTFISTIGIDFKIRTVELDGKKIKLQIWDTA 60
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APPLICANT: Pia M. Challita-Eid
APPLICANT: Pia M. Challita-Eid
APPLICANT: Arthur B. Ratiano
APPLICANT: Steve Chappell Mitchell
APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: Daniel E.H. Afar
APPLICANT: Aya Jakobovite
ITILE OF INVENTION: GTP-BINDING PROTEIN USEFUL IN TREATMENT
ITILE OF INVENTION: GTP-BINDING PROTEIN USEFUL
CURRENT APPLICATION NUMBER: US/09/834,765
CURRENT FILLING DATE: 2001-09-21
FRIOR APPLICATION NUMBER: 60/197,647
FRIOR APPLICATION NUMBER: 60/197,647
NUMBER OF SEQ ID NOS: 770
SOFTWARE: FastSEQ for Windows Version 4.0
SSEQ ID NO 766
SEQ ID NO 766
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 162;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 190;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
46.3%; Score 512; DB 10;
Best Local Similarity 47.7%; Pred. No. 1.6e-41;
Matches 93; Conservative 52; Mismatches 42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 766, Application US/09834765
Patent No. US20020055478A1
GENERAL INFORMATION:
                                                Sequence 5, Application US/09822860 Patent No. US20020146795A1
                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
CRGANISM: Discopyge ommata
US-09-822-860-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              186 NELALAELEEEEGKP 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  173 NENSLOEAVDKLKSP 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
COGANISM: Homo sapiens
US-09-834-765-766
                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -09-834-765-766
                                                                                                                                                                                                                                                                                                                                                        LENGTH: 190
                            09-822-860-5
                                                                                                                                                                                                                                                                                                                                SEQ ID NO 5
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                                                   RYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIGNKADEEQK 128
                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 MAKQYDVLFRLLLIGDSGVGKTCLLCRFTDNEFHSSHISTIGVDFKMKTIEVDGIKVRIQ 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIDLE
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ FOr Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/967,736
FILING DATE: 28-SEP-2001
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          43.3%; Score 478.5; DB 10;
45.0%; Pred. No. 2.7e-38;
tive 42; Mismatches 64; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Dr.
                                                                                                                                                         RQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRL 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: PF-0367 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 09/154,602
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                    COTLEY, Neil C. Shah, Purvi TITLE OF INVENTION: RAB PROTEINS NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-845-4166
                                                                                                                                                                                                                                                    US-09-967-736-3
Sequence 3, Application US/09967736
Patent No. US20020103340A1
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 201 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
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CLONE: 2514506
                                                                                                                                                                                                                                                                                                                                                          Lal, Preeti
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Best Local Similarity 45.09
Matches 91; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 94304
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ORGANISM: Homo sapien
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Best Local Similarity
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Best Local Similarity
Matches 84; Conserv
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US-09-794-257-16
                                                                                                                                                                                                                                                                                                             US-09-820-003A-4
                    RESULT 13
US-09-820-003A-4
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                                                                                                                                                                                                                                          SEQ ID NO 4
                                                                                                                                                                                                                                                                                 TYPE: PRT
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121 NKADEEQKRQVGREQGQQLAKEYGMDFYETSACTNLNIKESFTRLTELVLQAHRKELEGL 180
                    61 IWDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILLIG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 201;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFFWARE: FastSEO for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/967,736
FILING ATE: 28-Sep-2001
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    42.4%; Score 469; DB 10; 51.2%; Pred. No. 2.1e-37;
                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Incyte Pharmaceuticals, Inc. STREET: 3174 Porter Dr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: PF-0367 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 8:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    36; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 09/154,602
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE DESCRIPTION: SEQ ID NO: 8:
                                                                                                                                                                                                                                             ial, Preeti
Corley, Neil C.
Shah, Purvi
TITLE OF INVENTION: RAB PROTEINS
                                                                                                                                                                        Sequence 8, Application US/09967736
Patent No. US20020103340A1
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
                                                              181 RMRASNELALAELEEEGKPEG 202
                                                                                      176 GAASGERPNLKIDSTPVKPAG 197
                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    83; Conservative
                                                                                                                                                                                                                                                                                                                                                                          CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 94304
                                                                                                                                                           US-09-967-736-8
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08-08-967-136-8
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Matches 8
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Sequence 16, Application US/09794257
Patent No. US20020009804A1
GENERAL INFORMATION:
APPLICANT: Meyers, Rachel
TITLE OF INVENTION: 32705, 23224, 27423, 32700, 32712, No. US20020009804A1e1
TITLE OF INVENTION: HUman G-Proteins
FILE REPERENCE: 35800/2802485
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US/09/794,257
PRIOR APPLICATION NUMBER: 60/185,606
PRIOR FILING DATE: 2000-02-29
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 16
LENGTH: 198
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                                          GENERAL INFORMATION:
APPLICANT: MERKULOV, GENDADA Et al.
APPLICANT: MERKULOV, GENDADA ET AL.
TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
TITLE OF INVENTION: PROTEINS, AND USES THEREOF
FILE REFERENCE: CLO01196
CURRENT FILING DATE: 2001-03-29
NUMBER OF SEQ ID NOS: 39
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 IMDTAGQERYQTITKQYYRRAQGIFLVYDISSERSYQHIMKWVSDVDEYAPEGVQKILIG 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            141 NKCDLTTKKVVDYTTAKEFADSLGIPFLETSAKNATNVEQSF 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 10;
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Pred. No. 3.8e-37;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Pfam accession number PF00071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34; Mismatches
Sequence 4, Application US/09820003A Patent No. US20020142382A1
                                                                                                                                                                                                                                                                                                                                                                                                                                     42.3%; 51.9%; 1
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US-09-945-173-5

Sequence 5, Application US/09945173

Sequence 5, Application US/09945173

Patent No. US20020127568A1

GENERAL INFORMATION:

APPLICANT: MILLennium Pharmaceuticals, Inc.

APPLICANT: Meyers, Rachel

TITLE OF INVENTION: 47324, A NOVEL HUMAN G-PROTEIN AND USES

TITLE OF INVENTION: 47324, A NOVEL HUMAN G-PROTEIN AND USES

FILE REFERENCE: 38155-20035.00

CURRENT APPLICATION NUMBER: US/09/945,173

CURRENT FILING DATE: 2000-09-01

PRIOR FILING DATE: 2000-09-01

NUMBER OF SEQ ID NOS: 11

SOSTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 5

LENGTH: 198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 39.3%; Score 434.5; DB 10; Length 198; Best Local Similarity 50.9%; Pred. No. 3.9e-34; Matches 89; Conservative 30; Mismatches 43; Indels 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTHER INFORMATION: Consensus amino acid sequence US-09-945-173-5
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Search completed: January 16, 2003, 04:26:09 Job time : 78 secs

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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/916,901
FILING DATE: Filed Herewith
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
US-09-075-454-8

US-09-075-454-11

US-08-773-421-6

US-08-773-423-6

US-08-773-423-6

US-08-773-421-6

US-08-774-411-6

US-08-774-946A-5

PCT-US95-06420-5

US-08-842-3068-1

US-08-842-3068-4

US-08-973-1

US-08-973-1

US-09-078-317-1

US-09-078-317-1

US-09-078-317-1

US-09-078-317-1

US-09-078-317-1

US-09-078-317-1

US-09-156-424-1

US-09-156-428-1

US-09-387-341-1

US-09-388-483-1

US-09-488-1
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US-09-387-341-150
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REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0367 ITELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
TELEFAX: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: RAB PROTEINS NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-916-901-4
; Sequence 4, Application US/08916901
; Patent No. 5892012
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APPLICANT: Hillman, Jennifer
APPLICANT: Lal, Preeti
APPLICANT: Corley, Neil C.
APPLICANT: Shah, Purvi
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CITY: Palo Alto
STATE: CA
COUNTRY: USA
      FILING DATE:
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-LOOPEXT=0 -UNITS=bits -START=1 -END=1 -MATRIX=blosum62 -TRANS=human40.cdi
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-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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6: /cgn2_6/ptodatra1/ina/PcTUS_COMB.seq:*
                              GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                             - nucleic search, using frame_plus_p2n model
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US-09-154-602-4
US-08-824-813-2
US-09-108-1184-2
US-09-075-454-10
US-09-499-913-66
US-08-888-077A-28
US-08-773-423-4
US-08-773-423-4
US-08-773-421-4
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Listing first 45 summaries
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Perfect score:
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<u>2</u> Result

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AsnLysAlaAspGluGluGlnLysArgGlnValGlyArgGluGlnGlnGlnLeuAla 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            181 ArgMetArgAlaSerAsnGluLeuAlaLeuAlaGluLeuGluGluGluGluGlyLysPro 200
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Mismatches:
Indels:
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                                                                                                                                            Length:
Matches:
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APPLICANT: Lal, Preeti
APPLICANT: Corley, Neil C.
                                                                                                                                          2.39e-51
478.50
65.84%
45.05%
43.30%
      SEQUENCE CHARACTERISTICS:
ERNGTH: 925 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
LIBRARY: LIVERUT04
CLONE: 2514506
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                      Percent Similarity:
Best Local Similarity:
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US-09-154-602-4
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Mismatches:
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                                                                                                                                                                 SOFTWARE: FastSEQ CON Windows Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/154,602 FILING DATE:
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Matches:
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                                             ADDRESSEE: Incyte Pharmaceuticals,
                                                                                                                                                                                                                                                                                                PF-0367
         TITLE OF INVENTION: RAB PROTEINS NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                        PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/916,901
                                                                                                                                                                                                                                                 FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF
                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                  TELERAX: 415-845-4166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
TENGTH: 925 base pairs
TYPE: nucleic acid
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478.50
65.84%
45.05%
43.30%
                                                                                                                                                  IBM Compatible
                                                         3174 Porter Dr
                                                                                                                                     Diskette
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MEDIUM TYPE: Diskett
Shah, Purvi
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CLONE: 2514506
                                                                                                                                                                                                                                                                                                                                                                                                             linear
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Best Local Similarity:
                                                                        Palo Alto
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                                                                                                USA
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                                                                                  GACTCTCTGGGCATCCCCTTCTTGGAGAGGGCCCAAGAATGCCACCAATGTCGAGCAG 545
                        141 LysGluTyrGlyMetAspPheTyrGluThrSerAlaCysThrAsnLeuAsnIleLysGlu 160
                                                                                                                               161 SerPheThrArgLeuThrGluLeuValLeuGlnAlaHisArgLysGluLeuGluGlyLeu 180
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Conservative:
Mismatches:
Indels:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
OPTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATE:
APPLICATION TO THE FILE FILED HERWITH APPLICATION 1435
FILING DATE: FILED HERWITH
CLASSIFICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
APPLICANT: Guegler, Karl
TITLE OF INVENTION: NOVEL RAB PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length:
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                                                                                                                                                                                                                                                                                                                                                                  Sequence 2, Application US/08824873
Patent No. 5843717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 1340 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5e-46
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Best Local Similarity:
Query Match:
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US-08-824-873-2
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441 TGGACTGCCCTTCATGGAGACCAGCGCCAAGACGGGCCTCAACGTGGACTTGGCCTTCAC 500
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                                      25 Leu-CysArgPheThrAspAsnGluPhe---HisSerSerHisIleSerThrIleGlyVa 43
                                                                                                                                                                                                       43 lAspPheLysMetLysThrIleGluValAspGlyIleLysValArgIleGlnIleTrpAs 63
                                                                                                                                                                                                                                                                                        PThrAlaGlyGlnGluArgTyrGlnThrIleThrLysGlnTyrTyrArgArgAlaGlnGl 83
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SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
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3174 Porter Drive
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APPLICANT: Hillman, Jennifer L.
APPLICANT: Guegler, Karl
TITLE OF INVENTION: NOVEL RAB PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/198,184 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/09198184 Patent No. 6010859
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NAME: Billings, Lucy J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Palo Alto
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Lal, Preeti

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APPLICANT:
APPLICANT:
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                Length:
Matches:
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         REFERENCE/DOCKET NUMBER: PF-0240 US TELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-0555 TELEPHONE: 415-845-4166 INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: LENGTH: 1340 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 10, Application US/09075454 Patent No. 6391580 GENERAL INFORMATION: APPLICANT: Hillman, Jennifer L. APPLICANT: Tang, Y. Tom
36,749
                                                                                                                                                                                                                                           6.5e-46
437.50
67.84%
45.23%
39.59%
                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: single
REGISTRATION NUMBER:
                                                                                                                                                          IMMEDIATE SOURCE:
LIBRARY: PANCNOT04
CLONE: 738957
                                                                                                                                                linear
                                                                                                                                                                                                                                                                             Percent Similarity:
Best Local Similarity:
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DB:
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Mismatches:
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                                                                                                                        3: INCYTE PHARMACEUTICALS, INC. 3174 Porter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                            PF-0168-1 CIP
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                                                                                                                                                                                                                                                                        SOFTWARE: Word Perfect 6.1/MS-DOS 6.2
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/075.454
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                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA
APPLICATION NUMBER: 08/766,551
FILING DATE: DECEMBER 12, 1996
ATTORNEY/AGENT INFORMATION:
                                APPLICANT: Patterson, Chandra
APPLICANT: Batra, Sajeev
APPLICANT: Baughn, Mariah R.
TITLE OF INVENTION: RAS PROTEINS
NUMBER OF SEQUENCES: 14
Guegler, Karl J.
Corley, Neil C.
Patterson, Chandra
Batra, Sajeev
                                                                                                                                                                                                                                                                                                                                                                                                                                          39,132
                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Cerrone, Michael C. REGISTRATION NUMBER: 39,132 REFERENCE/DOCKET NUMBER: PF TELECOMMUNICATION INFORMATION: 650-855-0555
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429.00
68.48%
44.57%
38.82%
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MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
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LENGTH: 875 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                Herewith
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                                                                                                               CORRESPONDENCE ADDRESS:
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CLONE: 1528559
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Query Match:
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APPLICANT: Betty, Maria
APPLICANT: Betty, Maria
APPLICANT: Betty, Maria
APPLICANT: Lidg, Huai-Ping
APPLICANT: Lidg, Huai-Ping
APPLICANT: An, Wengian
TITLE OF INVENTION: POTASSIUM CHANNEL INTERACTORS AND USES THEREFOR
FILE REFERENCE: MI-070CP2
CURRENT APPLICATION NUMBER: US/09/399,913
CURRENT FILING DATE: 1999-09-21
EARLIER PILING DATE: 1998-11-30
EARLIER FILING DATE: 1998-11-25
EARLIER FILING DATE: 1998-11-25
EARLIER FILING DATE: 1999-11-20
EARLIER FILING DATE: 1999-04-23
EARLIER FILING DATE: 1999-04-23
EARLIER FILING DATE: 1999-07-09
EARLIER FILING DATE: 1999-07-09
EARLIER FILING DATE: 1999-07-09
EARLIER FILING DATE: 1999-07-09
SARLIER FILING DATE: 1999-07-09
SARLIER FILING DATE: 1999-07-09
NUMBER OF SEQ ID NOS: 73
COFTWARE PREAINT OF US OF THE CONTROL OF THE 
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TIGCTICTGCTGTATGACATCACAAATCTTCTTTCGACAACATCAGGGCCTGGCTC 390
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                                                                                                  SerAspValAspGluTyrAlaProGluGlyValGlnLysIleLeuIleGlyAsnLysAla
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Matches:
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Best Local Similarity:
Query Match:
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LOCATION: (1)
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Pred. No.:
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US-09-399-913-66
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312 TTGCTTCTGCTGTATGACATCACCAACAAATCTTTTGACAACAACAACAACGGCCTGGCTC 371
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TACGACCTCACGGGCAAGGTGATGCTTCTGGGAGACACAGGCGTCGGCAAAAAAATGTTTC 150
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; OTHER INFORMATION: Incyte ID No. 6426186 412477.1CB1
US-09-484-970B-142
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Matches:
Conservative:
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Sequence 142, Application US/09484970B
Sequence 142, Application US/09484970B
Sequence 142, Application US/09484970B
SEQUENCE INFORMATION:
APPLICANT: Volkmuth, Wayne
APPLICANT: Walker, Michael G
TITLE OF INVENTION: BONE REMODELING GENES
FILE REFERENCE: PB-0014 US
CURRENT APPLICATION UNMBER: US/09/484,970B
CURRENT FILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 172
SOFTWARE: PERL PROGRAM
SOFTWARE: PERL PROGRAM
SEQ ID NO 142
LENGTH: 2612
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44.57%
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ORGANISM: Homo sapiens
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Query Match:
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                                                                                                                                                                                                                                                      105 AspvalAspGluTyrAlaProGluGlyValGlnLysIleLeuIleGlyAsnLysAlaAsp 124
                                                                                                                                                                                                                                                                                        366
                                                                                                                                                                                                                                                                                                                    125 GluGluGlnLysArgGlnValGlyArgGluGlnGlnGlnGeuAlaLysGluTyrGly 144
                                                                                                                                                                                                                                                                                                                                        164
                                                                                                                                                                                                                                                                                                                                                                                                                                                165 LeuThrGluLeuVal-LeuGlnAlaHisArgLySGluLeuGluGlyLeuArgMetArg-- 183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              487 ACAGCAAAAGAAATTTATGAAAAATCCAAGAAGGGGTCTTTGACATTAATAATGAGGCA 546
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               184 -AlaserAsnGluLeuAlaLeuAlaGluLeu-------------GluGluGl 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   :::::::::||||||||||:::
547 AACGGCATCAAAATTGGCCCTCAGCATGCTACCAATGCATCTCACGGAGGAACCAA 606
                                                                                                                                                                                       PheLeuValTyrAspIleSerSerGluArgSerTyrGlnHisIleMetLysTrpValSer 104
                                                                                                                                                                                                             67 IIGCTACAGITIACAGACAAGAGGITICAGCCGGIGCAIGACCTCACAAIIGGIGIAGAG 126
                                                                            PheLysMetLysThrIleGluValAspGlyIleLysValArgIleGlnIleTrpAspThr 64
                                                                                                                         AlaGlyGlnGluArgTyrGlnThrIleThrLysGlnTyrTyrArgArgAlaGlnGlyIle 84
                                                                                                                                                 LeuCysArgPheThrAspAsnGluPheHisSerSerHisIleSerThrIleGlyValAsp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 28, Application US/0888077A
Patent No. 6020143
Patent No. 6020143
Patent INFORMATION:
GEORGE-HYSLOP, PETER H
APPLICANT: ST. GEORGE-HYSLOP, PETER H
APPLICANT: FRANENS, JOHANNA M
APPLICANT: FRANEN, PAUL E
TITLE OF INVENTION: GENETIC SEQUENCES AND PROTEINS RELATED
TITLE OF INVENTION: TO ALZHEIMER'S DISEASE AND USES THEREFOR.
NUMBER OF SEQUENCES: 41
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: LERNER, DAVID, LITTENBERG, KRUMHOLZ & MENTLIK
STREET: 600 SOUTH AVENUE WEST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/888,077A
FILING DATE: 03-UL-1997
CLASSIFICATION: 530
PRIOR APPLICATION: 530
APPLICATION NUMBER: US 08/592,541
FILING DATE: 26-JAN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              196 uGluGlyLysProGluGlyProAla 204
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REGISTRATION NUMBER: 36,629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 07090-1497
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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STATE: NJ
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163 ThrargLeuThrGluLeuValLeuGlnAlaHisArgLysGluLeuGluGlyLeuArg--- 181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              124 AspGluGluGlnLysArgGlnValGlyArgGluGlnGlnGlnGlnLeuAlaLysGluTyr 143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            104 SerAspValAspGluTyrAlaProGluGlyValGlnLysIleLeuIleGlyAsnLysAla 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84 IlePheLeuValTyrAspIleSerSerGluArgSerTyrGlnHisIleMetLysTrpVal 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              112 GCCTTATTGGTTTATGACATTGCTAAACATCTCACATATGAAATGTAGAAGGGTGGCGTG 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          44 AspPheLysMetLysThrIleGluValAspGlyIleLysValArgIleGlnIleTrpAsp 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ThralaglyGlnGluArgTyrGlnThrIleThrLysGlnTyrTyrArgArgAlaGlnGly 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 GlnTyrAspValLeuPheArgLeuLeuLeulleGlyAspSerGlyValGlyLysThrCys 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        432 GAICTACGTCATCTCAGGGCAGTTCCTACAGATGAAGCAAGAGCTTTTGCAGAAAAAAT
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SCHERING 3.0-017 CIP CIP
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Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-817-198A-2 (1-212) x US-08-888-077A-28 (1-970)
                                                                                                                                                                                                                                                                                                                    Length:
Matches:
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                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: 1..970
OTHER INFORMATION: /note= "Y2H9"
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REFERENCE/DOCKET NUMBER: SC
TELECOMMUNICATION INFORMATION
TELEPHONE: (908) 654-5000
                                                   INFORMATION FOR SEQ ID NO: 28
SEQUENCE CHARACTERISTICS:
LENGTH: 970 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                      7.94e-43
                                                                                                                                                                                                                                                                                                                                     411.50
64.04%
41.87%
37.24%
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Best Local Similarity:
Query Match:
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                                                                                                                                                                                                                                                              US-08-888-077A-28
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123 AlaAspGluGluGlnLysArgGlnValGlyArgGluGlnGlyGlnGlnLeuAlaLysGlu 142
                                                                                                                                                                                                    509 AATGGACTGCTCTTCCTGGAGACCTCAGCCCTGGACTCTACCAATGTTGAGCTAGCCTTT 568
                                                                                                                                                                                                                                                                                                                         569 GAGACTGTCCTGAAAGAAATCTTTGCGAAGGTGTCCAAGCAGAGACAGAACAGCATCCGG 628
                                                                                                                                                                                                                                                                                                                                                                   182 MetArgAlaSerAsnGluLeuAlaLeuAlaGluLeuGluGluGluGluGlglglyLysProGlu 201
                                                                                                                                                                                                                                                                                                                                                                                                       -----ACCAATGCCATCACTCTGGGC----AGTGCCCAGGNTGGACAGGAGCCT 673
                                         103 ValSerAspValAspGluTyrAlaProGluGlyValGlnLysIleLeuIleGlyAsnLys 122
                                                                                                                                                               449 AGTGACCTCAGCCAGGGCCGGGAAGTGCCCACTGAGGAGGCCCGAATGTTCGCTGAAAAC 508
                                                                                                                                                                                                                                                                                    ---ThrArgLeuThrGluLeuValLeuGlnAlaHisArgLysGluLeuGluGlyLeuArg 181
  329 GGGGCCCTCCTGGTGTTTGACCTAACCAAGCACCAGACCTATGCTGTGGTGGAGCGATGG 388
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Bandman, Olga
APPLICANT: Bandman, Olga
APPLICANT: Au-Young, Janice
TITLE OF INVENTION: NOVEL RAB PROTEINS
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
STREET: 3174 Porter Drive
CITY: Palo Alto
STATE: CA
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        674 GGCCCTGGGGAGAGAGGGCCTGTTGC 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                 202 GlyProAlaAsnSerSerLysThrCys 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: DOS
SOSTWARE: FASTEDO VERSION 1.5
SOSTWENT APPLICATION DATA:
APPLICATION NUMBER: US/08/741,411
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ATTORNEY/AGENT INFORMATION:
NAME: B1111095, Lucy J.
REGISTRATION NUMBER: 36, 749
REFERENCE/DOCKET NUMBER: PF-0.
TELECOMMUNICATION INFORMATION:
"PT-RPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-741-411-4
; Sequence 4, Application US/08741411
; Patent No. 6124116
; GENERAL INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
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LENGTH: 890 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLONE: Consensus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GlyilePheLeuValTyrAspIleSerSerGluArgSerTyrGlnHisIleMetLysTrp 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         269 GACACAGCTGGCCTGGAGCGGTACCGAGCCATCACCTCGGCGTACTATCGTGGTGCAGTG 328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 LysGlnTyrAspValLeuPheArgLeuLeuLeulleGlyAspSerGlyValGlyLysThr 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ValAspPheLysMetLysThrIleGluValAspGlyIleLysValArgIleGlnIleTrp 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    63 AspThrAlaGlyGlnGluArgTyrGlnThrIleThrLysGlnTyrTyrArgArgAlaGln 82
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                                                                                                                                    OPERATION SYSTEM: DOS SOFTWARE: FASTSEQ for Windows Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/773,423 FILING DATE: Herewith CLASSIFICATION: 514 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-817-198A-2 (1-212) x US-08-773-423-4 (1-847)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length:
Matches:
                                                                                                                                                      ADDRESSEE: Incyte Pharmaceuticals, Inc.
STREET: 3174 Porter Drive
CITY: Palo Alto
                                   APPLICANT: Hillman, Jennifer L.
APPLICANT: Goli, Surya K.
APPLICANT: Bandman, Olga
TITLE OF INVENTION: NOVEL RAB PROTEINS
NUMBER OF SEQUENCES: 9
ADDRESSPONDENCE ADDRESS:
ADDRESSEE: Incyte Pharmaceuticals, In
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: PF-0183 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Billings, Lucy J. REGISTRATION NUMBER: 36,749
                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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59.818
36.848
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SEQUENCE CHARACTERISTICS:
LENGTH: 847 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Percent Similarity:
Best Local Similarity:
Query Match:
                       GENERAL INFORMATION:
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                                                                                                                                                                                                                                            USA
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TOPOLOGY: lin
Patent No. 5869291
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                           94304
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                                                                                                                                                                                                                       STATE: C. COUNTRY:
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Palo Alto

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123 AlaAspGluGluLysArgGlnValGlyArgGluGlnGlyGlnGlnLeuAlaLysGlu 142
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132 TACGICCACCAGCTTCTCCCAGCACTACCGGGCCACCATCGGGGACTTCGCCCTC 191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          162 PheThrArgLeuThrGluLeuValLeuGlnAlaHisArgLysGluLeuGluGlyLeuArg 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      182 MetArgAlaSerAsnGluLeuAlaLeuAlaGluLeuGluGluGluGluGlyLySProGlu 201
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                                                                                                                                                                                                                                                                                                     67 GlnGluArgTyrGlnThrIleThrLysGlnTyrTyrArgArgAlaGlnGlyIlePheLeu
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                                           Conservative:
                                                          Mismatches:
Indels:
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                                                                                                                         US-09-817-198A-2 (1-212) x US-08-741-411-4 (1-890)
           Length:
Matches:
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Patent No. 6391580
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer L.
                         343.00
343.00
57.89%
36.36%
31.04%
                                                         Best Local Similarity:
Query Match:
                                           Percent Similarity
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US-09-075-454-13
Alignment Scores:
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Mismatches:
Indels:
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Matches:
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                                                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/766,551
FILING DATE: DECEMBER 12, 1996
ATTORNEY/AGENT INFORMATION:
NAME: CETROR, MICHAEL C.
REGISTRATION NUMBER: 39,132
REFERENCE/DOCKET NUMBER: PF-0168-
                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 650-855-0555
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340.50
60.31%
38.66%
30.81%
                                                                   IBM Compatible
                                                                                                                                                                                                                                                                                                                                         LENGTH: 803 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: Herewith
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                                                      Diskette
                                        COMPUTER READABLE FORM MEDIUM TYPE: Diskett
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SOFTWARE: Word Pe
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Best Local Similarity:
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CLONE: 2703745
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IMMEDIATE SOURCE
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              COUNTRY:
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123 AlaAspGluGluGlnLysArgGlnValGlyArgGluGlnGly
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APPLICANT: Batra, Sajeev
TYLICANT: Baughn, Mariah R.
TITLE OF INVENTION: RAS PROTEINS
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/766,551
FILING DATE: DECEMBER 12, 1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 8, Application US/09075454 Patent No. 6391580 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hillman, Jennifer L.
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TELEPHONE: 650-855-0555
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Corley, Neil C.
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CITY: Palo Alto
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            IBM Compatible
                                                                                                                                                                                                                                                                                            198 GlyLysPro------
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Lal, Preeti
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MEDIUM TYPE: Diskette
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APPLICANT: Knuth, Alexander
APPLICANT: Knuth, Alexander
APPLICANT: Char, Yao-Tseng
APPLICANT: Chen, Yao-Tseng
TITLE OF INVENTION: Cancer Associated Antigen Encoding Nucleic Acid Molecules
TITLE OF INVENTION: and Uses Thereof
FILE REFERENCE: LUD-5638
CURRENT APPLICATION NUMBER: US/09/493,914
CURRENT FILING DATE: 2000-01-28
NUMBER OF SEQ ID NOS: 5
                                                                                             111 :: 346 TCTGCACCTTGCTCGAGAAGTGAAAAGAGAACTGACAGGATGACAGGATCCATTAC 605
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142 uTyrGlyMetAspPheTyrGluThrSerAlaCysThrAsnLeuAsnIleLysGluSerPh 162
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Matches:
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Patent No. 6448073
                                                                                                                                                                                                                                                                                          APPLICANT: Jger, Dick
APPLICANT: Stockert, Elizabeth
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321.50
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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LENGTH: 1407
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DB:
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140 AlaLysGluTyrGly----MetAspPheTyrGluThrSerAlaCysThrAsnLeuAsnIle 158
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--GlnGlnLeu 139
                                                    432 TGTGAC-----CAGGGGAAGGATGTGCTCATGACAATGGCCTCAAGATGGACCAGTTC 485
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REFERENCE/DOCKET NUMBER: PF-0168-1 CIP
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/075,454
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Mismatches:
Indels:
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                                                                               Length:
Matches:
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APPLICANT: Corley, Neil C.
APPLICANT: Patterson, Chandra
APPLICANT: Batra, Sajeev
APPLICANT: Batra, Sajeev
TITLE OF INVENTION: RAS PROTEINS
NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11, Application US/09075454 Patent No. 6391580 GENERAL INFORMATION:
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Tang, Y. Tom
Lal, Preeti
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          LIBRARY: KIDNNOTO5
CLONE: 627565
                                                                                                   Percent Similarity:
Best Local Similarity:
Query Match:
IMMEDIATE SOURCE:
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STREET: 31
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APPLICANT:
                                       US-09-075-454-8
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Mismatches:
Indels:
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Matches:
                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 39,132
REFERENCE/DOCKET NUMBER: PF-0168-1 CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-855-0555
                                                                                                                                              6.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps:
                                                                                                                                        SOFTWARE: Word Perfect 6.1/MS-DOS CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/075,454
                                                                                                                                                                                                                                                         APPLICATION NUMBER: 08/766,551
FILING DATE: DECEMBER 12, 1996
ATTORNEY/AGENT INFORMATION:
NAME: Cerrone, Michael C.
REGISTRATION NUMBER: 39,132
                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 11:
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                                                                                       MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
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LENGTH: 1533 base pairs
                                                                                                                                                                                                     Herewith
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                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                        COMPUTER READABLE FORM:
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Palo Alto
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Best Local Similarity:
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US-09-075-454-11
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                                                                                                                                                                                                         FILING DATE:
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                                   COUNTRY:
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1045
                                    965 TACTGGGCAGTCTCATCTCTCACTGGTGAGAATGTCCGAGAATTCTTCTTCTTCCGTGTGGCA 1024
147 PheTyrGluThrSerAlaCysThrAsnLeuAsnIleLysGluSerPheThrArgLeuThr 166
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                                                                                                                                                                     187 GluLeuAlaLeuAlaGluLeuGluGluGluGluGlyLysProGluGly 202
                                                                                                                                                                                                                                                                                                           APPLICANT: Hillman, Jennifer L.
APPLICANT: Bandman, Olga
APPLICANT: Bandman, Olga
APPLICANT: Hawkins, Phillip R.
APPLICANT: Goli, Surya K.
TITLE OF INVENTION: NOVEL HUMAN GTP-BINDING PROTEINS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCTTE PHARMACEUTICALS, INC.
STREET: 3174 POrter Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length:
Matches:
Conservative:
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Indels:
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SOFTWARE: FastSEQ Version 2.0
CURENT APPLICATION DATA:
APPLICATION NUMBER: US/08/766,551
FILING DATE: Herewith
CLASSIPICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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ATORNEYAGENT INFORMATION:
NAME: B111ings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/POCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                      RESULT 15
US-08-766-551-6
; Sequence 6, Application US/08766551
; Patent No. 5840569
                                                                                               1025 GCACTGACCTTTGAGGCCAAT----
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LENGTH: 1255 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: SEQ ID NO:6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94304
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
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627051
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STATE: CA
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US-08-766-551-6
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                                                                                                      90 IleSerSerGluArgSerTyrGlnHisIleMetLysTrpValSerAsp----ValAspGlu 108
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                                                                                                                                                                                                                                                                                                                                                                                                                                   882 GCTCAGTATGCGCTGATGGAGAAGAMGCCCTCCAGGTGGCCCAGGGGATGAAGGCTGAG 941
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                                                                                                                                                                                                               702 TICAAAIGCATIGCATCAACCIACIAIAAGAGGAGCICAAGCCAICAICAITGICITCAAC 761
522 AAGGTCATTGTGGGGGGACCTGTCGGTGGGGAAGACTTGCCTCATTAATAGGTTCTGC 581
                                   30 AspAsnGluPheHisSerSerHisIleSerThrIleGlyValAspPheLysMetLysThr 49
                                                                                                                                                                             187 GluLeuAlaLeuAlaGluLeuGluGluGluGluGlyLysProGluGly 202
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Job time : 52 secs
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Perfect score:

Sequence:

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Run on:

Scoring table:

Total number

Searched:

88

Minimum Maximum

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Sequence 88. Application US/09764868; Patent No. US20020168711A1.; GENERAL INFORMATION:
; TITLE OF INFORMATION:
; TITLE OF INFORMATION:
; FILE REFERENCE: PTZ32; CURRENT APPLICATION NUMBER: US/09/764,868; CURRENT FILIGA DATE: 2001-01-17; Prior application data removed - refer to PALM or file wrapper; NUMBER OF SEQ ID NOS: 1510; SOFTWARE: Patentin Ver. 2.0; SEQ ID NO 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to PALM or file wrapper
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10.05-09-918-08-1461

10.05-09-164-868-71

10.05-09-156-88-93

10.05-09-156-88-94

10.05-09-156-88-113

10.05-09-918-8-113

10.05-09-918-8-113

10.05-09-918-8-148-113

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Conservative:
Mismatches:
US-09-925-300-631
US-09-794-257-9
US-09-834-975-879
US-09-834-975-879
US-09-834-975-895
US-09-834-975-896
US-09-938-842A-836
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Best Local Similarity:
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-Q-CQG12_1/V2PTO_28001/US9817198 | Tunat_13012003_120052_22048 | Appp_query.fasta_1.391
-Q-CQG12_1/V2PTO_28001/US9817198 | Tunat_13012003_120052_22048 | Appp_query.fasta_1.391
-QDB-Published_Applications_NA -QFMT-fastap -SUFFIX-rnpb -MINMATCH-0.1
-LOOPCL-0 -LOOPEXT-0 -UNITS-bits -START-1 -END--1 -MATRIX-blosum62
-TRANS-human40.cdl -LIST-45 -DOCALIGN-200 -THR_SCORE-pct -THR_MAX-100
-TARANS-human40.cdl -LIST-45 -DOCALIGN-200 -NORM-ext -HEAPSIZE-500 -MINLEN-0
-MAXLEN-200000000 -USER-US09817198_GCGN_11_24_@runat_13012003_120052_22048
-NCPU-6 -ICPU-3 -NO.MAMP -LARGEQUERY -NG_SCORES-0 -WAIT -LONGLOG
-NCPU-120 -WARN_TIMEOUT-30 -THREADS-1 -XGAPOP-10 -XGAPEXT=0.5 -FGAPOP-6
-FGAPEXT-7 -YGAPOP-10 -YGAPEXT=0.5 -DELOP-6 -DELEXT=7
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Sequence 1, Appli
Sequence 507, App
Sequence 91, Appl
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1750.446 Million cell updates/sec
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Copyright (c) 1993 - 2003 Compugen Ltd
                                                               - nucleic search, using frame_plus_p2n model
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10 US-09-817-198A-1
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10 US-09-925-302-91
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Fatent No. US20020146758A1
GENERAL INFORMATION:
APPLICANT: YE, Jame et al.
TITLE OF INVENTION: ISOLATED HUMAN RAS-LIKE PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING THESE HUMAN RAS-LIKE
TITLE OF INVENTION: PROTEINS, AND USES THEREOF
FILE REFERENCE: CL001188
CURRENT APPLICATION NUMBER: US/09/817,198A
CURRENT FILING DATE: 2001-03-27
NUMBER OF SEQ ID NOS: 33
SOFTWARE: FatESQ for Windows Version 4.0
SEQ ID NOS: 30
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Query Match:
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TYPE: DNA
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APPLICANT: Rosen et al.

TITLE OF INVENTION: Nucleic Acids, Proteins, and Antiboo FILE REFERENCE: PT232
CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT FILING DATE: 2001-01-17
Prior application data removed - refer to PALM or file NUMBER OF SEQ ID NOS: 1510
SOFTWARE: Patentin Ver. 2.0
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OTHER INFORMATION: n ec
NAME/KEY: SITE
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OTHER INFORMATION: n ec
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OTHER INFORMATION:
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LOCATION: (493)
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; APPLICANT: Rosen et al.; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies; FILE REFERENCE: PA104; CURRENT APPLICATION NUMBER: US/09/925,302; CURRENT FILING DATE: 2001-08-10; PRIOR APPLICATION NUMBER: PCT/US00/05918; PRIOR PFLICATION NUMBER: 60/124,270; PRIOR FILING DATE: 1999-03-12; NUMBER OF SEQ ID NOS: 896; SOFTWARE: Patentin Ver. 2.0
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Mismatches:
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US-09-764-868-507
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APPLICANT: Steve Ruben
TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
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102
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Mismatches:
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Matches:
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; Patent No. US20020151681A1
; GENERAL INFORMATION:
                      a,t,g,
                                                                             a,t,g,
; LOCATION: (1264)
; OTHER INFORMATION: n equals a,
; NAME/KEY: misc feature
; LOCATION: (1268)
; OTHER INFORMATION: n equals a,
US-09-925-302-91
                                                                                                                                           8.4e-58
534.50
66.38%
43.40%
                                                                                                                                                                              Percent Similarity:
Best Local Similarity:
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                                                                                                                                Alignment Scores:
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FILE REFERENCE: 35800/209285
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                                                                                            LENGTH: 624
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Matches:
Conservative:
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   FILE REFERENCE: PA101
CURRENT APPLICATION WUMBER: US/09/925,300
CURRENT TLING DATE: 2001-08-10
CURRENT PILING DATE: 2001-08-10
PRIOR PILING DATE: 2000-03-08
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1890
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: 1537
                                                                                                                                                                                 COCATION: (5)
COTHER INFORMATION: n equals a,t,g, or US-09-925-300-631
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; Sequence 9, Application US/09794257
; Patent No. US20020009804A1
                                                                                                                                                                                                                                          3.5e-57
530.50
82.56%
55.81%
                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                       NAME/KEY: misc feature
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Best Local Similarity:
Query Match:
DB:
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APPLICANT: Meyers, Rachel
TITLE OF INVENTION: 32705, 23224, 27423, 32700, 32712, No. US20020009804A1e1
TITLE OF INVENTION: Human G-Proteins
TITLE OF INVENTION: Human G-Proteins
CURRENT APPLICATION NUMBER: US/09/794,257
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION UNDER: 60/185,606
NUMBER OF SEQ ID NOS: 16
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Conservative:
Mismatches:
Indels:
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CURRENT APPLICATION NUMBER: US/09/794,257
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: 60/185,606
PRIOR FILING DATE: 2000-02-29
NUMBER OF SEQ ID NOS: 16
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 7, Application US/09794257; Patent No. US20020009804A1; GENERAL INFORMATION:
                                                                                                                                                                                                                           1.71e-57
528.50
80.68%
53.98%
47.83%
                                                                                                                                        TYPE: DNA
CRGANISM: homo sapiens
US-09-794-257-9
                                                                                                                                                                                                                                                             Percent Similarity:
Best Local Similarity:
Query Match:
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NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
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                  SOFTWARE: Fas:
SEQ ID NO 879
LENGTH: 2497
                                                          TYPE: DNA
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APPLICANT: Lillie, James
APPLICANT: Brown, Jeffrey
APPLICANT: Brown, Jeffrey
APPLICANT: Born, Jeffrey
APPLICANT: Born, Jeffrey
APPLICANT: Wan Huffel, Christophe
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: OF HUMAN CANCERS
FILE REFERENCE: MRI-0168
CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
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                                                                                                                                                                                             Mismatches:
                                                                                                                                                     Length:
Matches:
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  for Windows Version 4.0
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Patent No. US20020110815A1
GENERAL INFORMATION:
                                                                                                                                            4.17e-57
528.50
80.68%
53.98%
47.83%
                        LENGTH: 1161
TYPE: DNA
ORGANISM: homo sapiens
                                                                                            ; LOCATION: (18)...(641)
US-09-794-257-7
                                                                                                                                                                            Percent Similarity:
Best Local Similarity:
SOFTWARE: FastSEQ
SEQ ID NO 7
                                                                                 NAME/KEY: CDS
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Sequence 885, Application US/09834975
Sequence 885, Application Sequence Applicant: Inlie, James
APPLICANT: Brown, Jeffers
APPLICANT: Brown, Jeffers
APPLICANT: Wan Huffel, Christophe
APPLICANT: Wan Huffel, Christophe
TITLE OF INVENTION: NOT THE IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY
TITLE OF INVENTION: OF HUMAN CANCERS
FILLE REPERENCE: MRI-0168
CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
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Matches:
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Windows Version 4.0
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528.50
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47.83%
                                                                                                                               NAME/KEY: misc_feature LOCATION: (1)...(2497)
                                                                                   ORGANISM: Homo sapiens
                                                                                                                                                       ; LOCATION: (1)...(2497)
; OTHER INFORMATION: n =
US-09-834-975-879
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APPLICANT: Van Huffel, Christophe
TITLE OF INVENTION: NOVEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY
TITLE OF INVENTION: OF HUMAN CANCERS
FILE REFERENCE: MRI-016B
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Mismatches:
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Matches:
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PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 1046
SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 885
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Patent No. US20020110815A1
GENERAL INFORMATION:
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NAME/KEY: misc_feature
LOCATION: (1)...(2497)
OTHER INFORMATION: n = A,T,C or
US-09-834-975-885
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80.68%
53.98%
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                                                                             TYPE: DNA
ORGANISM: Homo sapiens
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Query Match:
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Sequence 896, Application US/09834975
Sequence 896, Application US/09834975
Patent No. US20020110815A1
GENERAL INFORMATION:
APPLICANT: Lillie, James
APPLICANT: Bolt, Andrew
APPLICANT: Bolt, Andrew
APPLICANT: Wan Huffel, Christophe
TITLE OF INVENTION: NOYEL GENES, COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE IDENTIFICATION, ASSESSMENT, PAND THERAPY
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Mismatches:
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Matches:
CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 1046
SOFWMARE: FastSEQ for Windows Version 4.0
SOFWMARE: PastSEQ for Windows Version 4.0
LENGTH: 2497
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NAME/KEY: misc_feature
LOCATION: (1)...(2497)
CTHR INFORMATION: n = US-09-834-975-894
                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
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Best Local Similarity:
Query Match:
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US-09-834-975-896
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us-09-817-198a-2.rnpb

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TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING TITLE OF INVENTION: SAME, AND METHODS OF USE FILE REPERENCE: SCRIP1300-3 842A CURRENT APPLICATION NUMBER: US/09/938,842A CURRENT FILING DATE: 2001-08-24 PRIOR PLILING DATE: 2001-08-24 PRIOR PLILING DATE: 2000-08-24 PRIOR PLILING DATE: 2000-08-24 PRIOR FILING DATE: 2000-08-24 PRIOR FILING DATE: 2001-01-16 PRIOR FILING DATE: 2001-01-16 PRIOR FILING DATE: 2001-01-16 PRIOR FILING DATE: 2001-06-22 NUMBER OF SEQ ID NOS: 5379
SEQ ID NOS: 5379
SEQ ID NOS: 5379
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Mismatches:
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Best Local Similarity:
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Conservative:
Mismatches:
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FILE REFERENCE: MRI-016B
CURRENT APPLICATION NUMBER: US/09/834,975
CURRENT FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/197,538
PRIOR FILING DATE: 2000-04-14
NUMBER OF SEQ ID NOS: 1046
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 896
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                                                                                                                                                                                     NAME/KEY: misc_feature
LOCATION: (1)...(2497)
OTHER INFORMATION: n = A,T,C or
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528.50
80.68%
53.98%
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APPLICANT: Kreps, Joel
APPLICANT: Wang, Xun
APPLICANT: Zhu, Tong
                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                    Best Local Similarity:
Query Match:
                                                                                                                                                                                                                                                                                                        Percent Similarity:
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                                                                                                                                                                                                                                                                             Pred. No.:
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81 AlaGlnGlyIlePheLeuValTyrAspIleSerSerGluArgSerTyrGlnHisIleMet 100
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Best Local Similarity:
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91
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OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
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                                                                                                                                                      ADDRESSEE: Incyte Pharmaceuticals, Inc. STREET: 3174 Porter Dr. CITY: Palo Alto
                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/967,736
FILING DATE: Sep-2001
PRIOR APPLICATION NUMBER: US/09/967,736
FILING DATE: CURROWN:
APPLICATION NUMBER: 09/154,602
FILING DATE: CURROWN:
NAME: BILINGS, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0367 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-85-055
TELEPHONE: 415-845-4166
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
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Matches:
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Gaps:
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                                                                              Lal, Preeti
Corley, Neil C.
Shah, Purvi
TITLE OF INVENTION: RAB PROTEINS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
                      Sequence 4, Application US/09967736
Patent No. US20020103340A1
GENERAL INFORMATION:
APPLICANT: Hillman, Jennifer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 925 base pairs
                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
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478.50
65.84%
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CLONE: 2514506
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Best Local Similarity:
Query Match:
                                                                                                                                                                                                    STATE: CA
                                                                                                                                                                                                                  COUNTRY:
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           US-09-967-736-4
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161 SerPheThrArgLeuThrGluLeuValLeuGlnAlaHisArgLySGluLeuGluGlyLeu 180
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                                                                                                                                                                                                                                                                                                                                                                                                           591 GGAGCAGCCTCTGGGGGGGGAGCGGCCCAATCTCAAGATCGACAGCACCCCTGTAAAGCCG 650
                                                                                                                                                                                                          141 LysGluTyrGlyMetAspPheTyrGluThrSerAlaCysThrAsnLeuAsnIleLysGlu 160
                                                                                                                             121 AsnLysAlaAspGluGluGlnLysArgGlnValGlyArgGluGlnGlyGlnLouAla 140
                                                                                                                                                        486 GACTCTCTGGGCATCCCCTTCTTGGAGACGAGGCGCCAAGAATGCCAACAATGTCGAGCAG
101 LysTrpValSerAspValAspGluTyrAlaProGluGlyValGlnLysIleLeuIleGly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Davis, Keith R.
APPLICANT: Allon, Keith R.
APPLICANT: Hoffman, Neil
APPLICANT: Hoffman, Neil
APPLICANT: Hurban, Patrick
TITLE OF INVENTION: Expressed Sequences of Arabidopsis;
TITLE OF INVENTION: thaliana
FILE REFERENCE: 2023US (PARA-012PRV)
CURRENT FILING NATE: 2001-01-26
PRIOR APPLICATION NUMBER: US 60/178,472
PRIOR FILING DATE: 2001-01-26
PRIOR FILING DATE: 2001-01-27
PRIOR FILING DATE: 2000-01-27
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Mismatches:
Indels:
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Matches:
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SOFTWARE: FastSEQ for Windows Version 4.0
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Matthew, Abraham V.
Ledford, Brooke L.
Woessner, Jeffrey P.
Haas, William David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Arabidopsis thaliana
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An, Yong-Qiang
Hamilton, Carol M.
Price, Jennifer L.
Raines, Tracy M.
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477.50
65.578
45.288
43.218
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Rameaka, Joshua G.
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Kricker, Maja
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609 84 36 49 0

Conservative:

Mismatches: Length: Matches:

Indels:

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AlaGlnGlyIlePheLeuValTyrAspIleSerSerGluArgSerTyrGlnHisIleMet 100
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                                  6.59e-50
468.00
71.018
49.708
42.35$
                                                                   Percent Similarity:
Best Local Similarity:
 JS-09-938-842A-832
                            Alignment Scores:
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Sequence 80220020160378A1

GENERAL INFORMATION:

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Harper, Jeff

APPLICANT: Wang, Xun

APPLICANT: Wang, Xun

TITLE OF INVENTION: STRESS-REGULATED GENES OF PLANTS, TRANSGENIC PLANTS CONTAINING

TITLE OF INVENTION: SAME, AND METHODS OF USE

TITLE OF INVENTION: SAME, AND METHODS OF USE

FILE REFRENCE: SCRIP1300-3

CURRENT PAPLICATION NUMBER: US/09/938,842A

CURRENT FILING DATE: 2001-08-24

PRIOR APPLICATION NUMBER: US 60/227,866

PRIOR APPLICATION NUMBER: US 60/227,866

PRIOR PELING DATE: 2001-01-16

PRIOR FILING DATE: 2001-01-16

PRIOR FILING DATE: 2001-01-16

PRIOR FILING DATE: 2001-01-16

NUMBER OF SEQ ID NOS: 5379

SEQ ID NOS: 5379
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                                                                                                                                                                             AACAAGTCTGATCTTACTGAAAACAGAGCCATTCCTTATGAAACTGCCAAGGCTTTTGCC
                                                     MetalaLysGlnTyrAspValLeuPheArgLeuLeuLeuIleGlyAspSerGlyValGly
                                                                                                           LysThrCysLeuLeuCysArgPheThrAspAsnGluPheHisSerSerHisIleSerThr
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Gaps:
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TYPE: DNA
ORGANISM: Arabidopsis thaliana
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US-09-938-842A-832
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